

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization
International Bureau(43) International Publication Date
10 January 2002 (10.01.2002)

PCT

(10) International Publication Number
WO 02/02587 A1

- (51) International Patent Classification⁷: C07H 21/04, C12N 15/10, 15/11, 15/12
- (21) International Application Number: PCT/US01/20917
- (22) International Filing Date: 29 June 2001 (29.06.2001)
- (25) Filing Language: English
- (26) Publication Language: English
- (30) Priority Data:
60/215,135 30 June 2000 (30.06.2000) US
60/225,266 14 August 2000 (14.08.2000) US
- (71) Applicant (*for all designated States except US*): HUMAN GENOME SCIENCES, INC. [US/US]; 9410 Key West Avenue, Rockville, MD 20850 (US).
- (72) Inventors; and
- (75) Inventors/Applicants (*for US only*): FISCELLA, Michele [IT/US]; 6308 Redwing Road, Bethesda, MD 20817 (US). NI, Jian [CN/US]; 17815 Fair Lady Way, Germantown, MD 20874 (US). RUBEN, Steven, M. [US/US]; 18528 Heritage Hills Drive, Olney, MD 20832 (US).
- (74) Agents: HOOVER, Kenley et al.; 9410 Key West Avenue, Rockville, MD 20850 (US).
- (81) Designated States (*national*): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW.
- (84) Designated States (*regional*): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).
- Published:**
- with international search report
 - before the expiration of the time limit for amending the claims and to be republished in the event of receipt of amendments
 - with (an) indication(s) in relation to deposited biological material furnished under Rule 13bis separately from the description
 - with sequence listing part of description published separately in electronic form and available upon request from the International Bureau
- For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.*



WO 02/02587 A1

(54) Title: B7-LIKE POLYNUCLEOTIDES, POLYPEPTIDES, AND ANTIBODIES

(57) Abstract: The present invention relates to novel human B7-like polypeptides and isolated nucleic acids containing the coding regions of the genes encoding such polypeptides. Also provided are vectors, host cells, antibodies, and recombinant methods for producing human B7-like polypeptides. The invention further relates to diagnostic and therapeutic methods useful for diagnosing and treating disorders related to these novel human B7-like polypeptides.

INTERNATIONAL SEARCH REPORT

Int. application No.
PCT/US01/20917

A. CLASSIFICATION OF SUBJECT MATTER

IPC(7) : C07H 21/04; C12N 15/10, 15/11, 15/12

US CL : 536/23.1, 23.5; 435/69.1, 326, 320.1, 455

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 536/23.1, 23.5; 435/69.1, 326, 320.1, 455

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched
NONE

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

WEST, DIALOG, BIOSIS, CA, EMBASE, MEDLINE

search terms: fiscella, ni, ruben, b7, b7-1, b7-2, cd80, cd86

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	WO 00/36107 A (CORIXA CORPORATION) 22 JUNE 2000, see entire document, particularly SEQ ID NO: 391	1-10, 14, 15

☐ Further documents are listed in the continuation of Box C. ☐ See patent family annex.

* Special categories of cited documents:	"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
"A" document defining the general state of the art which is not considered to be of particular relevance	"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
"E" earlier document published on or after the international filing date	"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
"L" document which may throw doubt on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)	"A" document member of the same patent family
"O" document referring to an oral disclosure, use, exhibition or other means	
"P" document published prior to the international filing date but later than the priority date claimed	

Date of the actual completion of the international search 01 OCTOBER 2001	Date of mailing of the international search report 16 NOV 2001
Name and mailing address of the ISA/US Commissioner of Patents and Trademarks Box PCT Washington, D.C. 20231 Facsimile No. (703) 305-3330	Authorized officer PHILLIP GANIBEL Telephone No. (703) 308-0796

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US01/20917

Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This international report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:

2. ☐ Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:

3. ☐ Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

Please See Extra Sheet.

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:

4. ☒ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:
1-10, 14, 15

Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
☐ No protest accompanied the payment of additional search fees.

INTERNATIONAL SEARCH REPORT

In international application No.
PCT/US01/20917

BOX II. OBSERVATIONS WHERE UNITY OF INVENTION WAS LACKING
This ISA found multiple inventions as follows:

This application contains the following inventions or groups of inventions which are not so linked as to form a single inventive concept under PCT Rule 13.1. In order for all inventions to be searched, the appropriate additional search fees must be paid.

Groups 1-49, claims 1-10, 14, 15, all in part, drawn to an isolated nucleic acids of SEQ ID NO: X or encoding a peptide of SEQ ID NO: Y, wherein X and Y are values that correlates to those listed in Table 1 and correspond to one of the cDNA clone IDs, respectively as well as vectors host cells and methods of making a proteins.

For example, If Group 1 is elected, this correlates to Gene No. 1, ATCC Deposit No. PTA02332, SE ID NO: 2 and SEQ ID NO: Y

It is noted that the Groups would be numbering 7, if the X and Y sequences are limited to each row. The Groups number 49, if one can pick and choose a separate X and a separate Y from Table 1.

Applicant is invited to clarify the number of possibilities intended.

Groups 50-98, claims 11, 12 and 16, all in part, drawn to proteins comprising sequences encoded by SEQ ID NO: X and a peptide of SEQ ID NO: Y, wherein X and Y are values that correlates to those listed in Table 1 and correspond to one of the cDNA clone IDs, respectively.

Groups 99-147, claim 13, all in part, drawn to an antibody that binds a protein comprising sequences encoded by SEQ ID NO: X and a peptide of SEQ ID NO: Y, wherein X and Y are values that correlates to those listed in Table 1 and correspond to one of the cDNA clone IDs, respectively.

Groups 148-196, claim 17, all in part, drawn to methods of preventing or treating a medical conditions with an isolated nucleic acids of SEQ ID NO: X or encoding a peptide of SEQ ID NO: Y, wherein X and Y are values that correlates to those listed in Table 1 and correspond to one of the cDNA clone IDs, respectively.

Groups 197-245, claim 18, all in part, drawn to methods of diagnosing a pathological condition via an isolated nucleic acids of SEQ ID NO: X or encoding a peptide of SEQ ID NO: Y, wherein X and Y are values that correlates to those listed in Table 1 and correspond to one of the cDNA clone IDs, respectively.

Groups 246-294, claim 19, all in part, drawn to methods of diagnosing a pathological condition via an antibody that binds a protein encoded by isolated nucleic acids of SEQ ID NO: X or encoding a peptide of SEQ ID NO: Y, wherein X and Y are values that correlates to those listed in Table 1 and correspond to one of the cDNA clone IDs, respectively

Groups 295-343, claims 20-21, all in part, drawn to methods of identifying a binding partner of a peptide encoded by isolated nucleic acids of SEQ ID NO: X or encoding a peptide of SEQ ID NO: Y, wherein X and Y are values that correlates to those listed in Table 1 and correspond to one of the cDNA clone IDs, respectively

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US01/20917

Groups 344-392, claim 22, all in part, drawn to methods of preventing or treating a medical condition with a protein encoded by the nucleic acids of SEQ ID NO: X or encoding a peptide of SEQ ID NO: Y, wherein X and Y are values that correlates to those listed in Table 1 and correspond to one of the cDNA clone IDs, respectively.

This application contains claims directed to more than one species of the generic invention. These species are deemed to lack Unity of Invention because they are not so linked as to form a single inventive concept under PCT Rule 13.1. In order for more than one species to be searched, the appropriate additional search fees must be paid. The species are as follows:

The polynucleotides and polypeptides of each of the claims in Table 1 are unrelated, each to the other. The polynucleotides sequence encode structurally distinct polypeptides and do not share a special technical feature. Further the technical feature that links the DNA, proteins, antibody and methods is not a contribution over the prior art of Corixa Corporation (WO 00/36107), particularly SEQ ID NO: 391. Also, see the Search Report. Thus, the technical feature of the polynucleotide sequence is not special and the Groups are not so linked under PCT Rule 13.1. Additionally the claimed methods encompassed different ingredients, process steps and endpoints, which are not so coextensive and which do not share the same technical feature.

The polynucleotides and polypeptides of each of the clones in Table 1 are unrelated, each to other. The polynucleotides sequences encode structurally distinct polypeptides and do not share a special technical feature. Furthermore, the technical feature that links the DNA, protein, antibody and methods of PTA-2332 is not a contribution over the prior art of Corixa Corporation (WO 00/36107), particularly SEQ ID NO: 391 set forth in the Search Report. Thus, the technical feature of the polynucleotide sequence is not special and the Groups are not so linked under PCT Rule 13.1. Additionally, the claimed methods encompass different ingredients, process steps and endpoints which are not coextensive and which do not share the same technical feature.

<110> Human Genome Sciences, Inc.

<120> B7-Like Polynucleotides, Polypeptides, and Antibodies

<130> PT124PCT

<140> Unassigned

<141> 2001-06-29

<150> 60/215,135

<151> 2000-06-30

<150> 60/225,266

<151> 2000-08-14

<160> 49

<170> PatentIn Ver. 2.0

<210> 1

<211> 733

<212> DNA

<213> Homo sapiens

<400> 1

gggatccgga	gcccaaatct	tctgacaaaa	ctcacacatg	cccaccgtgc	ccagcacctg	60
aattcgaggg	tgacaccgtca	gtcttcctct	tcccccaaa	acccaaggac	accctcatga	120
tctcccgga	tcctgaggtc	acatgcgtgg	tggtggacgt	aagccacgaa	gacctgagg	180
tcaagttcaa	ctggtacgtg	gacggcgtgg	aggtgcataa	tgccaagaca	aagccgcggg	240
aggagcagta	caacagcacg	taccgtgtgg	tcagcgtcct	caccgtcctg	caccaggact	300
ggctgaatgg	caaggagtac	aagtgcagg	tctccaacaa	agccctccca	accccatcg	360
agaaaaccat	ctccaaagcc	aaagggcagc	cccagaaacc	acaggtgtac	accctgcccc	420
catcccgga	tgagctgacc	aagaaccagg	tcagcctgac	ctgcctgggc	aaaggcttct	480
atccaagcga	catcgccgtg	gagtgggaga	gcaatgggca	gccggagaac	aactacaaga	540
ccacgcctcc	cgtgctggac	tccgacggct	ccttcttctc	ctacagcaag	ctcaccgtgg	600
acaagagcag	gtggcagcag	gggaacgtct	tctcatgtct	cgtgatgcat	gaggctctgc	660
acaaccacta	cacgcagaag	agcctctccc	tgtctccggg	taaatgagtg	cgacggccgc	720
gactctagag	gat					733

<210> 2

<211> 3357

<212> DNA

<213> Homo sapiens

<400> 2

caccagcagt	agtagcagaa	gcgaagagcg	caaacgcaac	cgctctcccc	gcgcgttggc	60
cgattcatta	atgcagctgg	cacgacaggt	ttcccactg	gaaagcgggc	agtgagcgca	120
acgcaattaa	tgtgagttag	ctcactcatt	aggcacccca	ggctttacac	tttatgcttc	180
cggtctgcat	gttgtgtgga	attgtgagcg	gataacaatt	tcacacagga	aacagctatg	240
accatgatta	cgccaagctc	gaaattaacc	ctcactaaag	ggaacaaaag	ctggagctcc	300
accgcggtgg	cgcccgctct	agaactagt	gatcccccg	gctgcaggaa	ttcggcacga	360
gaggcagcgg	cagctccact	cagccagtac	ccagatacgc	tggaacctt	ccccagccat	420
ggcttccctg	gggcagatcc	tcttctggag	cataattagc	atcatcatta	ttctggctgg	480
agcaattgca	ctcatcattg	gctttggtat	ttcaggggaga	cactccatca	cagtactac	540
tgtcgctca	gctgggaaca	ttggggagga	tggaatcctg	agctgcactt	ttgaacctga	600
catcaaacct	tctgatatcg	tgatacaatg	gctgaaggaa	ggtgttttag	gcttgggtcca	660
tgagttcaaa	gaaggcaaag	atgagctgtc	ggagcaggat	gaaatgttca	gaggccggac	720
agcagtgttt	gctgatcaag	tgatagtgtg	caatgcctct	ttgcggctga	aaaacgtgca	780

actcacagat	gctggcacct	acaaatgtta	tatcatcact	tctaaaggca	aggggaatgc	840
taaccttgag	tataaaactg	gagccttcag	catgccggaa	gtgaatgtgg	actataatgc	900
cagctcagag	accttgcggt	gtgaggctcc	ccgatgggtc	ccccagccca	cagtggctctg	960
ggcatcccaa	gttgaccagg	gagccaactt	ctcgggaagtc	tccaatacca	gctttgagct	1020
gaactctgag	aatgtgacca	tgaaggttgt	gtctgtgctc	tacaatgtta	cgatcaacaa	1080
cacatactcc	tgtatgattg	aaaatgacat	tgccaaagca	acaggggata	tcaaagtgc	1140
agaatcgag	atcaaaaggc	ggagtcacct	acagctgcta	aactcaaagg	cttctctgtg	1200
tgtctcttct	ttctttgcc	tcagctgggc	acttctgcct	ctcagccctt	acctgatgct	1260
aaaataatgt	gccttggcca	caaaaaagca	tgcaaagtca	ttgttacaac	agggatctac	1320
agaactat	caccaccaga	tatgacctag	ttttatat	ctgggaggaa	atgaattcat	1380
atctagaagt	ctggagtgag	caaacaagag	caagaaacaa	aaagaaggca	aaagcagaag	1440
gtccaatat	gaacaagata	aatctatctt	caaagacata	ttagaagttg	ggaaaataat	1500
tcattgtgaac	tagacaagt	tggttaagag	gataagtaaa	atgcacgtgg	agacaagtgc	1560
atccccagat	ctcagggacc	tccccctgcc	tgtcacctgg	ggagtgagag	gacaggatag	1620
tgcatgttct	ttgtctctga	attttttagt	atatgtgctg	taatgttgct	ctgaggaagc	1680
ccctggaaag	tctatcccaa	catatccaca	tcttatattc	cacaaattaa	gctgtagtat	1740
gtaccctaag	acgctgctaa	tcgactgcc	cttcgcaact	caggggcggc	tgcattttag	1800
taatgggtca	aatgattcac	tttttatgat	gcttccaaag	gtgccttggc	ttctcttccc	1860
aactgacaaa	tgccaaagtt	gagaaaaatg	atcataat	tagcataaac	agagcagtcg	1920
gcgacaccga	ttttataaat	aaactgagca	ccttcttttt	aaacaaacaa	atgcgggttt	1980
atttctcaga	tgatgttcac	ccgtgaatgg	tccagggaag	gacctttcac	cttgactata	2040
tggcattatg	tcatacacaag	ctctgaggct	tctcctttcc	atcctgcgtg	gacagctaag	2100
acctcagttt	tcaatagcat	ctagagcagt	gggactcagc	tggggtgatt	tcgcccccca	2160
tctccggggg	aatgtctgaa	gacaattttg	gttacctcaa	tgagggagtg	gaggaggata	2220
cagtgtctact	accaactagt	ggataaaggc	cagggatgct	gctcaacctc	ctaccatgta	2280
caggacgtct	ccccattaca	actacccaat	ccgaagtgtc	aactgtgtca	ggactaagaa	2340
accctggttt	tgagtagaaa	agggcctgga	aagaggggag	ccaacaaatc	tgtctgcttc	2400
ctcacattag	tcattggcaa	ataagcattc	tgtctctttg	gctgctgcct	cagcacagag	2460
agccagaact	ctatcgggca	ccaggataac	atctctcagt	gaacagagtt	gacaaggcct	2520
atgggaaatg	cctgatggga	ttatcttcag	cttggtgagc	ttctaagttt	ctttcccttc	2580
attctaccct	gcaagccaag	ttctgtaaga	gaaatgcctg	agttctagct	caggttttct	2640
tactctgaat	ttagatctcc	agacccttcc	tggccacaat	tcaaattaag	gcaacaaaca	2700
tataccttcc	atgaagcaca	cacagacttt	tgaagcaag	gacaatgact	gcttgaattg	2760
aggccttgag	gaatgaagct	ttgaaggaaa	agaatacttt	gtttccagcc	cccttcccac	2820
actcttcatg	tgtaaccac	tgcttctctg	gaccttgag	ccacgggtgac	tgtattacat	2880
gttggttatag	aaaactgatt	ttagagttct	gatcgttcaa	gagaatgatt	aaatatacat	2940
ttcctaataa	aaaaaaaaaa	aaactcgagg	gggggcccgg	tacccaattc	gccctatagt	3000
gagtcgtatt	acaattcact	ggccgtcggt	ttacaacgtc	gtgactggga	aaaccctggc	3060
gttaccacaac	ttaatcgctt	tgacgcacat	ccccctttcg	ccagctggcg	taatagcgaa	3120
gaggcccgca	ccgatcgccc	ttcccaacak	ttgcgcagcc	tgaatggcga	atggcaaatt	3180
gtaagcggtta	atattttggt	aaaattcgcg	ttaaattttt	gttaaatacag	ctcatttttt	3240
aaccaatagg	ccgaaatcgg	caaaatccct	tataaatcaa	aagaatagac	cgagataggg	3300
ttgagtgttg	ttccagtttg	gaacaagagt	ccactattaa	agtgttcacc	gcggtga	3357

<210> 3

<211> 2406

<212> DNA

<213> Homo sapiens

<400> 3

ccacgcgtcc	ggaatgaaca	acttttcttc	tcttgaatat	atcttaacgc	caaattttga	60
gtgctttttt	gttaccatc	ctcatatgtc	ccagctggaa	agaatcctgg	gttgagcta	120
ctgcatgttg	attgttttgt	ttttcctttt	ggctgttcat	tttggtggct	actataagga	180
aatctaacac	aaacagcaac	tgttttttgt	tgtttacttt	tgcatcttta	cttgaggagc	240
tgtggcaagt	cctcatatca	aatacagaac	atgatcttcc	tcctgcta	gttgagcctg	300
gaattgcagc	ttcaccagat	agcagcttta	ttcacagtga	cagtccttaa	ggaactgtac	360
ataatagagc	atggcagcaa	tgtgacctg	gaatgcaact	ttgacactgg	aagtcagtgtg	420
aaccttgag	caataacagc	cagtttgcaa	aaggtggaaa	atgatacatc	cccacaccgt	480
gaaagacca	ctttgctgga	ggagcagctg	cccctaggga	aggcctcggt	ccacatacct	540
caagtcacaag	tgagggacga	aggacagtag	caatgcataa	tcattctatgg	ggcgccctgg	600

gactacaagt	acctgactct	gaaagtcaaa	gcttcctaca	ggaaaataaa	cactcacatc	660
ctaaagggttc	cagaaacaga	tgaggtagag	ctcacctgcc	aggctacagg	ttatcctctg	720
gcagaagtat	cctggccaaa	cgtcagcggt	cctgccaaaca	ccagccactc	caggaccctc	780
gaaggcctct	accagggtcac	cagtgttctg	cgccctaaagc	caccccctgg	cagaaacttc	840
agctgtgtgt	tctggaatac	tcacgtgagg	gaacttactt	tggccagcat	tgaccttcaa	900
agtcagatgg	aacccaggac	ccatccaact	tggctgcttc	acattttcat	cccctcctgc	960
atcattgctt	tcattttcat	agccacagt	atagccctaa	gaaaacaact	ctgtcaaaag	1020
ctgtattctt	caaaagacac	aacaaaaaga	cctgtcacca	caacaaagag	ggaagtgaac	1080
agtgtctgtg	atctgaacct	gtggtcttgg	gagccagggg	gacctgatat	gacatctaaa	1140
gaagcttctg	gactctgaac	aagaattcgg	tggcctgcag	agcttgccat	ttgcactttt	1200
caaatgcctt	tggatgacct	agcactttaa	tctgaaacct	gcaacaagac	tagccaacac	1260
ctggccatga	aacttgcccc	ttcactgate	tggactcacc	tctggagcct	atggctttaa	1320
gcaagcacta	ctgcacttta	cagaattacc	ccactggatc	ctggaccac	agaattcctt	1380
caggatcctt	cttgcctgcca	gactgaaagc	aaaagggaatt	atttcccctc	aagttttcta	1440
agtgtatttc	aaaagcagag	gtgtgtggaa	atttccagta	acagaaacag	atgggttgcc	1500
aatagagtta	ttttttatct	atagcttctt	ctgggtacta	gaagaggcta	ttgagactat	1560
gagctcacag	acagggcttc	gcacaaactc	aatcataat	tgacatgttt	tatggattac	1620
tggaatcttg	atagcataat	gaagtgttcc	taattaacag	agagcattta	aatatacact	1680
aagtgcacaa	attgtggagt	aaagtcatca	agctctgttt	ttgaggtcta	agtcacaaag	1740
catttggtttt	aacctgtaat	ggcaccatgt	ttaatgggtg	tttttttttt	gaactacatc	1800
tttcccttaa	aaattattgg	tttcttttta	tttgttttta	ccttagaaat	caattatata	1860
cagtcaaaaa	tatttgatat	gctcatacgt	tgtatctgca	gcaatttcag	ataagtagct	1920
aaaatggcca	aagcccaaaa	ctaagcctcc	ttttctggcc	ctcaatatga	ctttaaattt	1980
gacttttctg	tgctcagtt	tgacatctg	taatacagca	atgctaagta	gtcaaggcct	2040
ttgataattg	gcaactatgga	aatcctgcaa	gatccacta	catatgtgtg	gagcagaagg	2100
gtaactcggc	tacagtaaca	gcttaatttt	gttaaatgtg	ttctttatag	tggagccatg	2160
aagctcagag	cattagctga	cccttgaact	attcaaatgg	gcacattagc	tagtataaca	2220
gacttacata	ggtgggccta	agcaagctc	cttaactgag	caaaatttgg	ggcttatgag	2280
aatgaaaggg	tgtgaaattg	actaacagac	aatcatata	tctcagtttc	tcaattctca	2340
tgtaaatcag	agaatgcctt	taaagaataa	aactcaattg	ttattcttca	aaaaaaaaaa	2400
aaaaaa						2406

<210> 4

<211> 3059

<212> DNA

<213> Homo sapiens

<400> 4

ggcacgagct	gtcatccggt	tccatgccgt	gaggtccatt	cacagaacac	atccatggct	60
ctcatgctca	gtttggttct	gagtcctctc	aagctgggat	cagggcagtg	gcaggtgttt	120
gggacagaca	agcctgtcca	ggccttggtg	ggggaggacg	cagcattctc	ctgtttcctg	180
tctcctaaga	ccaatgcaga	ggccatggaa	gtgcggttct	tcaggggcca	gttctctagc	240
gtggtccacc	tctacagggg	cggaaggac	cagccattta	tgcatatgcc	acagtatcaa	300
ggcaggacaa	aactggtgaa	ggattctatt	gaggaggggc	gcatctctct	gaggctggaa	360
aacattactg	tgttggtatg	tggcctctat	gggtgcagga	ttagttccca	gtcttactac	420
cagaaggcca	tctgggagct	acaggtgtca	gcactgggct	cagttcctct	catttccatc	480
gcgggatatg	ttgatagaga	catccagcta	ctctgtcagt	cctcgggctg	gttcccccg	540
cccacagcga	agtggaaagg	tccacaagga	caggatttgt	ccacagactc	caggacaaac	600
agagacatgc	atggcctggt	tgatgtggag	atctctctga	ccgtccaaga	gaacgccggg	660
agcatatcct	gttccatgcg	gcatgctcat	ctgagccgag	aggtggaatc	cagggtacag	720
ataggagact	ggagaagaaa	gcacggacag	gcaggtaaaa	gaaaatatc	ctcttcacac	780
atttatgact	cctttccaag	tctctcggtt	atggattttt	atatcctgag	gcccgtgggt	840
cctgcagag	ccaagcttgt	gatgggaact	ctgaaattgc	agattctggg	ggaggtgcat	900
ttttagagag	agcccatag	ccttcttcag	atctctggag	ggtccacaac	actcaaaaag	960
ggtcccaatc	cttgggtctt	cccttctccc	tgcgcctgt	ttcccacgtg	agcacggaac	1020
tgctgtctct	ctctgcttgc	tttcagaatt	gagagacgcc	cggaaacacg	caggtacca	1080
cgctgagag	ggttaacagt	ggcctggagt	aggaagatga	ccagtacag	atatggagcc	1140
catccagctt	gtagacagca	aatctgtgat	gcccgaatcc	acccagggg	gcagctgcct	1200
ctaaatacac	ttcttggtcc	aggacttgga	gggaaaagcg	tagggactgg	gtcagctagg	1260
aggggtcaca	ggcaagacgc	caggggaactg	agggcattag	tagctggctt	ctaggggtct	1320

gtgcaaaggg	gaacgaagt	aagttagcag	gaactgggtg	gtggaaggaa	gctgaatcct	1380
ggagtcactc	aaggtctcac	aaagtc aaat	agagggctta	cgtgggaggg	cagtggtagg	1440
gctgggtgaa	catctcatgg	ttgagcatct	ccaagcatca	gtgaggcacg	ggggctgccc	1500
tggagaaggt	acatggctgg	tgggatagtg	ggactggccg	gacccctacc	ggagccagtc	1560
tgcagtggga	gggtcgacct	cttgctccag	cccagatttc	gtcttcagta	actcatgctt	1620
cctctctccc	ccaccgcacc	ccagtggagg	tgactctgga	tccagagacg	gctcaccgga	1680
agctctgcgt	ttctgatctg	aaaactgtaa	cccatagaaa	agctcctcag	gaggtgcctc	1740
actctgagaa	gagatttaca	aggaagagtg	tggtggcttc	tcagggtttc	caagcagggg	1800
aacattactg	ggaggtggac	gtgggacaaa	atgtaggggtg	gtatgtggga	gtgtgtcggg	1860
atgacgtaga	cagggggaag	aacaatgtga	ctttgtctcc	caacaatggg	tattgggtcc	1920
tcagactgac	aacagaacat	ttgtatttca	cattcaatcc	ccattttatc	agcctcccc	1980
ccagcacc	tcctacacga	gtaggggtct	tcctggacta	tgagggtggg	accatctcct	2040
tcctcaatac	aaatgaccag	tcctttat	ataccctgct	gacatgtcag	tttgaaggct	2100
tggtgagacc	ctatatccag	catgcgatgt	atgacgagga	aaaggggact	cccatattca	2160
tatgtccagt	gtcctgggga	tgagacagag	aagaccctgc	ttaaagggcc	ccacaccaca	2220
gacccagaca	cagccaaggg	agagtgtctc	cgacaggtgg	ccccagcttc	ctctccggag	2280
cctgcgcaca	gagagtcacg	ccccccactc	tcctttaggg	agctgagggt	cttctgccct	2340
gagccctgca	gcagcggcag	tcacagcttc	cagatgaggg	gggattggcc	tgaccctgtg	2400
ggagtcagaa	gccatggctg	ccctgaagtg	gggacggaat	agactcacat	taggtttagt	2460
ttgtgaaaa	tcctccagc	taagcgatct	tgaacaagtc	acaacctccc	aggctcctca	2520
tttctagtc	acggacagtg	attcctgcct	cacaggtgaa	gattaaagag	acaacgaatg	2580
tgaatcatgc	ttgcaggttt	gagggccaca	gtgtttgcta	atggatgtgt	ttttatgatt	2640
atacattttc	cccaccataa	aactctgttt	gccttaattc	ccacattaat	ttactttttc	2700
ctcctatacc	caaatccacc	catggaatag	tttaattggaa	cacctgcctt	tgtgaggctc	2760
caaagaataa	agaggaggtg	ggatttttca	ctgattctat	aagcccagca	ttacctgata	2820
ccaaaaccag	gcaaagaaaa	cagaagaaga	ggaaggaaaa	ctacaggtcc	atatccctca	2880
ttaacacaga	cacaaaaatt	ctaaataaaa	ttttaacaaa	ttaaactaaa	caatatat	2940
aaagatgata	tataactact	cagtgtggtt	tgtcccacaa	atgcagagtt	ggtttaatat	3000
ttaaatatca	accagtgtaa	ttcagcacat	taataaagta	aaaaaaaaaa	aaaaaaaaaa	3059

<210> 5

<211> 2682

<212> DNA

<213> Homo sapiens

<220>

<221> SITE

<222> (1)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (2)

<223> n equals a,t,g, or c

<400> 5

nncacgagcc	tgtgcccctg	gaaaggttgg	agacttgggg	gacgactgga	gaattgccat	60
ttgaggacca	aaggagaaaa	gaaactacac	gctaattcta	gaaggcctcc	tgtccctgcc	120
tgctctgggt	gctcatggaa	ccagctgctg	ccctgcactt	ctcccggcca	gcctccctcc	180
tcctcctcct	cagcctgtgt	gcactgggtc	cagcccagtt	tactgtcgtg	gggccagcta	240
atcccctcct	ggccatgggtg	ggagaaaaaca	ctacattacg	ctgccatctg	tcaccgaga	300
aaaatgctga	ggacatggag	gtgcggtggt	tccggtctca	gttctcccc	gcagtgtttg	360
tgtataaggg	tgggagagag	agaacagagg	agcagatgga	ggagtaccgg	ggaagaatca	420
cctttgtgag	caaagacatc	aacaggggca	gcgtggccct	ggtcatacat	aacgtcacag	480
cccaggagaa	tgggatctac	cgctgttact	tccaagaagg	caggtcctac	gatgaggcca	540
tcctacgcct	cgtggtggca	ggccttgggt	ctaagccctt	cattgaaatc	aaggcccaag	600
aggatggggg	catctggctg	gagtgcata	ctggaggggtg	gtaccagag	cccctcacag	660
tgtggaggga	cccctacggg	gaggttgtgc	cgcctctgaa	ggagggttcc	atcgctgatg	720
ctgacggcct	cttcatgggtc	accacagctg	tgatcatcag	agacaagtat	gtgagggaatg	780
tgtcctgctc	tgtcaacaac	accctgctcg	gccaggagaa	ggaaactgtc	atttttat	840

cagaatcctt	tatgccagc	gcatctccct	ggatggtggc	cctagctgtc	atcctgaccg	900
catctccctg	gatggtgtcc	atgactgtca	tcctggctgt	tttcatcadc	ttcatggctg	960
tcagcatctg	ttgcatcaag	aaacttcaaa	gggaaaaaaa	gattctgtca	ggggaaaaga	1020
aagttgaaca	agaggaaaaa	gaaattgcac	agcaacttca	agaagaattg	cgatggagaa	1080
gaacattctt	acatgctgct	gatgtggtcc	tggatccaga	caccgctcat	cccgagctct	1140
tcctgtcaga	ggaccggaga	agtgtgaggc	ggggccccta	caggcagaga	gtgcctgaca	1200
acccagagag	attcgacagt	cagccttggt	tcctgggatg	ggagagcttc	gcctcagggg	1260
aacattacag	gggaaacttc	acagagtggg	gaccaccag	agcctataga	atcaattcct	1320
tggactcaca	gccatgcaga	aagccctggc	catctcagca	gccaccgcac	aaccccccta	1380
atgaaagaca	cgccctcctc	ccctctggtc	acgtaagaga	acatcttcca	gctgcctttt	1440
tcacaccac	tccagccctc	tgccccagtt	ttctcctcct	cactagtctg	tggctttagt	1500
agttcctttg	cttgtaatta	tgggatggga	tccaggcata	gggaactagt	tgtttcatag	1560
ctcccagtc	aaaagaaagt	gagagaagct	gttgggcagc	gaacctactg	tttaaaatca	1620
ggataaccac	attaagccca	atatgccagt	tggcaccaga	tgctgtggac	ttggaatgag	1680
gccaacaggg	ttcaccagga	tgagagagga	gagaggaatc	cacaggacca	ccagaagggg	1740
gagggaaaca	gatatgcaga	tcagagatag	aggaagtgtt	gagaggaaag	gggaggtcct	1800
gctgattcct	cagaatggct	tctggaccct	ggagatgttt	ggaaaccaat	accgggccct	1860
gtcctccctt	gagaggattc	tccttttgaa	ggagtccctt	tgccgggtgg	gcgtcttctt	1920
ggactatgaa	gctggagatg	tctccttcta	caacatgagg	gacagatcac	acatctacac	1980
atgtcccctg	tcagccttta	atgtgcctgt	gaggccattc	ttcagggttag	ggtctgatga	2040
cagccccatc	ttcatctgcc	ctgcactcac	aggagccagt	ggggtcatgg	tgctgaaga	2100
gggcctgaaa	cttcacagag	tggggaccca	ccaaggttgt	aaggatggct	aagtcccacc	2160
ataagagcta	aagggtcctg	ggagatgatg	gctcatttcc	acccaacccc	aggatttcca	2220
cagcacacac	ccacaggcct	ggacctggga	tgaagatgaa	tgaagaacat	ggactcatgt	2280
ggatgtggtt	tggctcagat	gtccctgcaa	taaacaaggg	gtcagtactt	agtccctgag	2340
tgtggttgag	gtttgaggtc	ctgggtcgagc	agggcagtac	tggaccaggt	ctacgtcagc	2400
attcaggttc	aatggggaca	ccagtggctt	caaacttcc	gatctaatta	tgtttttaga	2460
cacttagaag	ttattgagga	ctttaaagaa	cttttgttta	tttgggttaa	tatttatgac	2520
atttgaccat	tgaacaaaa	atttaaaatg	ttatctttta	atttatgtta	aaatagcatt	2580
aataaatcag	ttataggtta	atgtagatag	gatgttttgt	gaaaaagcaa	tctattgtgt	2640
ccaaataaaa	aaacaaaaag	tgtaaaaaaa	aaaaaaaaaa	aa		2682

<210> 6

<211> 1726

<212> DNA

<213> Homo sapiens

<220>

<221> SITE

<222> (1)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (2)

<223> n equals a,t,g, or c

<400> 6

nmcgattcgg	ctccaaactc	cggcgctgca	gccgatcgga	ctctggggccg	cggtgggcac	60
cgcgcgagc	tagggagccg	agaaccgcgg	cgagcccga	ggacggccag	agcgcgaggg	120
tcgctgcgc	tcgcagagcc	ggagccgagt	cgagccgggc	gcccgggctg	cctggagacg	180
ccgtgacttt	gaagtgtaac	ttcaagacag	atgggcgcac	gcgggagatc	gtgtggtacc	240
gggtgacgga	tgttggcacc	atcaagcaaa	agatcttcac	cttcgacgcc	atgttctcca	300
ccaactactc	acacatggag	aactaccgca	agcgagagga	cctggtgtac	cagtccactg	360
tgaggtgccc	cgaggtccgg	atctcagaca	atggtcccta	tgagtgccat	gtgggcatct	420
acgaccgcgc	caccagggag	aaggtggtcc	tggcatcagg	caacatcttc	ctcaacgtca	480
tggctcctcc	cacctccatt	gaagtgggtg	ctgctgacac	accagcccc	tcagccgct	540
accaagccca	gaacttcacg	ctggtctgca	tcgtgtctgg	aggaaaacca	gcacccatgg	600
tttatttcaa	acgagatggg	gaaccaatcg	acgcagtgcc	cctatcagag	ccaccagctg	660
cgagctccgg	ccccctacag	gacagcaggc	ccttccgcag	ccttctgcac	cgtgacctgg	720

atgacaccaa	gatgcagaag	tcaactgtccc	tcctggacgc	cgagaaccgg	ggtgggcgac	780
cctacacgga	gcgccctcc	cgtggcctga	ccccagatcc	caacatcctc	ctccagccaa	840
ccacagagaa	cataccagag	acggtcgtga	gccgtgagtt	tccccgctgg	gtccacagcg	900
ccgagcccac	ctacttcctg	cgccacagcc	gcaccccgag	cagtgcggc	actgtggaag	960
tacgtgccct	gtcacctgg	accctcaacc	cacagatcga	caacgaggcc	ctcttcagct	1020
gcgaggtcaa	gcacccagct	ctgtcgatgc	ccatgcaggc	agaggtcacg	ctggttgccc	1080
ccaaaggacc	caaaatttgt	atgacgcccc	gcagagcccg	ggtaggggac	acagtgagga	1140
ttctggtcca	tgggtttcag	aacgaagtct	tcccgagacc	catgttcacg	tggacgcggg	1200
ttgggagccg	cctcctggac	ggcagcgctg	agttcgacgg	gaaggagctg	gtgctggagc	1260
gggttcccg	cgagctcaat	ggctccatgt	atcgctgcac	cgcccagaac	ccactgggct	1320
ccaccgacac	gcacaccg	ctcatcgtgt	ttgaaaaccc	aaatatccca	agaggacgg	1380
aggactctaa	tggttccatt	ggccccactg	gtgcccggct	caccttggtg	ctcgccctga	1440
cagtgattct	ggagctgacg	tgaaggcacc	cgccccggcc	actccatcag	gcactgacat	1500
ctccgcgacc	ggttttcatt	tcttttctaa	actatttcca	gtcttggtct	tagtctcttt	1560
ccatctgtgt	cttggcttct	tcagtcggtt	taattaaaac	aaacagaaca	attttcccca	1620
caaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	1680
aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaa		1726

<210> 7

<211> 1021

<212> DNA

<213> Homo sapiens

<220>

<221> SITE

<222> (1)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (2)

<223> n equals a,t,g, or c

<400> 7

nncacgagcc	tgtgcccctg	gaaagggttg	agacttgagg	gacgactgga	gaattgccat	60
ttgaggacca	aaggagaaaa	gaaactacac	gctaattcta	gaaggcctcc	tgtccctgcc	120
tgtctctgggt	gctcatggaa	ccagctgctg	ccctgcactt	ctcccgccca	gcctccctcc	180
tcctcctcct	cagcctgtgt	gcactgggtct	cagcccagggt	cactgtcgtg	gggcccactg	240
atcccacctc	ggccatgggtg	ggagaaaaaca	ctacgttacg	atgctgtctg	tcacccgagg	300
aaaatgctga	ggacatggag	gtgcgggtggt	tccagtctca	gttctccctc	gcagtgtttg	360
tgtataagggt	tggagagag	agaacagagg	agcagaagga	ggagtaccga	gggagaacca	420
cctttgtgag	caaagacagc	aggggcagcg	tggccctgat	catacacaat	gtcacagccg	480
aggataacgg	catctaccag	tgttacttcc	aagaaggcag	gtcctgcaat	gaggccatcc	540
tgcaccttgt	ggtggcagac	cagcacaatc	ctctttcctg	gatccccatt	ccgcaggggg	600
cactctccct	atgaaaagaa	gattccaggg	gaaaaatcct	tcctcctgca	caagggccac	660
catgagttag	tttgccctgc	taagccgtgg	gcttgacttc	ttgagaagca	catgcagaac	720
tcagttgagg	ccatgagccg	ggggaaaatg	gtgaatctcg	gaagagaagt	cctatgcctg	780
ccttagcact	gagctgtgca	cttctgagag	tgagaggaga	caccatcaat	aattgtcttg	840
ggacaactgg	aataaacagt	gactgccag	agaactacga	tatttgaaat	cttatttctt	900
gatgaatatt	catcctgact	tctttcctga	aatgctgttt	gcaaagagag	tgacttatat	960
gtaagtagag	cgttttatta	aagcaagact	taatacagaa	gcaaaaaaaaa	aaaaaaaaaa	1020
a						1021

<210> 8

<211> 1835

<212> DNA

<213> Homo sapiens

<220>

<221> SITE
 <222> (1)
 <223> n equals a,t,g, or c

<220>
 <221> SITE
 <222> (2)
 <223> n equals a,t,g, or c

<400> 8
 nnacatccat ggctctaattg ctcaagctg ggtgggggag gacgcagcat 60
 agtggcagggt gtttgggcca gacaagcctg tccaggcctt ggtgggggag gacgcagcat 120
 tctctgtttt cctgtctcct aagaccaatg cagaggccat ggaagtgcgg ttcttcaggg 180
 gccagttctc tagcgtgggc cactcttaca gggacgggaa ggaccagcca tttatgcaga 240
 tgccacagta tcaaggcagg acaaaactgg tgaaggattc tattgcggag gggcgcatct 300
 ctctgaggct ggaaaacatt actgtgttgg atgtggcct ctatgggtgc aggtattgtt 360
 cccagtctta ctaccagaag gccatctggg agctacagggt gtcagcactg ggctcagttc 420
 ctctcatttc catcacggga tatgttgata gagacatcca gctactctgt cagtccctcg 480
 gctgggtccc cgggccca gcgaagtggg aaggtccaca aggacaggat ttgtccacag 540
 actccaggac aaacagagac atgcatggcc tggttgatgt ggagatctct ctgaccgtcc 600
 aagagaaacgc cgggagcata tctgttcca tgcggcatgc tcatctgagc cgagaggtgg 660
 aatccagggt acagatagga gatacctttt tgcagcctat atcgtggcac ctggctacca 720
 aagtactggg aatactctgc tgtggcctat ttttggcat tgttgactg aagattttct 780
 tctccaaatt ccagtggaaa atccaggcgg aactggactg gagaagaaag cacggacagg 840
 cagaattgag agacgcccgg aaacacgcag tggaggtgac tctggatcca gagacggctc 900
 accgaagct ctgcgtttct gatctgaaaa ctgtaacca tagaaaagct cccagggagg 960
 tgctcactc tgagaagaga tttacaagga agagtgtggg ggcttctcag agtttccaag 1020
 cagggaaca ttaactgggag gtggacggag gacacaataa aaggtggcgc gtgggagtgt 1080
 gccgggatga tgtggacagg aggaaggagt acgtgacttt gtctcccgat catgggtact 1140
 gggctcctcag actgaatgga gaacatttgt atttcacatt aaatccccgt tttatcagcg 1200
 tcttccccag gacccacct acaaaaatag gggctctcct ggactatgag tgtgggacca 1260
 tctcttctt caacataaat gaccagtccc ttatttatac cctgacatgt cggtttgaag 1320
 gcttattgag gccctacatt gagtatccgt cctataatga gcaaaatgga actcccagag 1380
 acaagcaaca gtgagtctc ctacacaggc accacgcct tcttccccag gggtgaaatg 1440
 taggatgaat cacatccac attcttctt agggatatta aggtctctct cccagatcca 1500
 aagtcgccga gcagccggcc aaggtggctt ccagatgaag ggggactggc ctgtccacat 1560
 gggagtccag tgtcatggct gccctgagct gggaggggag aaggctgaca ttacatttag 1620
 tttgctctca ctccatctgg ctaagtgatc ttgaaatacc acctctcagg tgaagaaccg 1680
 tcaggaaatc ccatctcaca ggctgtgtg tagattaagt agacaaggaa tgtgaataat 1740
 gcttagatct tattgatgac agagtgtatc ctaatggtt gttcattata ttacactttc 1800
 agtaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaa 1835

<210> 9
 <211> 2626
 <212> DNA
 <213> Homo sapiens

<400> 9
 aattcggcac gagaggcagc ggcagctcca ctcaagcagt acccaggata cgctggggaa 60
 ccttcccca gccatggctt ccctggggca gatcctcttc tggagcataa tttagcatca 120
 tcattattct ggctgaagca attgcactca tcattggctt tggattttca gggagacact 180
 ccatcacagt cactactgtc gcctcagctg ggaacattgg ggaggatgga atcctgagct 240
 gcacttttga acctgacatc aaactttctg atatcgtgat acaatggctg aaggaagggtg 300
 ttttaggctt ggtccatgag ttcaaagaag gccaaagatg agctgtcggg gcaggatgaa 360
 atgttcagag gccgggacag cagtgtttgc tgatcaagtg atagttggca atgcctcttt 420
 tgcggctgaa aaacgtgcaa ctacagatg ctggcaccta caaatgttat atcatcatt 480
 ctaaaaggcaa ggggaatgct aaccttgagt ataaaactgg agccttcagc atgccggaag 540
 tgaatgtgga ctataatgcc agctcagaga ccttgcggtg tgaggctccc cgatgggtcc 600
 cccagcccac agtgggtctgg gcatcccaag ttgaccaggg agccaacttc tcggaagtct 660
 ccaataccag ctttgagctg aactctgaga atgtgacct gaagggtgtg tctgtgctct 720

acaatgttac	gatcaacaac	acatactcct	gtatgattga	aatgacatt	gccaaagcaa	780
caggggatat	caaagtgaca	gaatcggaga	tcaaaaggcg	gagtcaccta	cagctgctaa	840
actcaaaggc	ttctctgtgt	gtctcttctt	tctttgccat	cagctgggca	cttctgcctc	900
tcagccctta	cctgatgcta	aaataatgtg	ccttggccac	aaaaaagcat	gcaaagtcac	960
tgttacaaca	gggatctaca	gaactatttc	accaccagat	atgacctagt	tttatatttc	1020
tgggaggaaa	tgaattcata	tctagaagtc	tggagtgagc	aaacaagagc	aagaaacaaa	1080
aagaagccaa	aagcagaagg	ctccaatatg	aacaagataa	atctatcttc	aaagacatat	1140
tagaagttgg	gaaaataatt	catgtgaact	agacaagtgt	gttaagagtg	ataagtaaaa	1200
tgcacgtgga	gacaagtgca	tccccagatc	tcagggacct	ccccctgcct	gtcacctggg	1260
gatgagagga	caggatagtg	catgttcttt	gtctctgaat	ttttagttat	atgtgctgta	1320
atgttgctct	gaggaagccc	ctggaaagtc	tatcccaaca	tatccacatc	ttatatccca	1380
caaattaagc	tgtagtatgt	accctaagac	gctgctaata	gactgccact	tcgcaactca	1440
ggggcggtcg	cattttagta	atgggtcaaa	cgattcagct	tttatgatgc	ttccaaaggt	1500
gccttggtct	ctcttcccaa	ctgacaaatg	ccaaaagttg	agaaaaatga	tcataatttt	1560
agcataaaca	gagcaagtcg	gcgacaccga	ttttataaat	aaactgagca	ccttcttttt	1620
aaacaaacaa	atgcgggttt	atctctcaga	tgatgttcat	cccgtgaatg	gtccagggaa	1680
ggacctttca	ccttgactat	atggcattat	gtcatcacia	gctctgaggc	ttctcctttc	1740
catectgctg	ggacagctaa	gacctcagtt	ttcaatagca	tctagagcag	tgggactcag	1800
ctggggtgat	ttcgcccccc	atctccgggg	gaatgtctga	agacaatttt	ggttacctca	1860
atgagggagt	ggaggaggat	acagtgtctac	taccaactag	tggataaagg	ccagggatgc	1920
tgctcaacc	tctaccatg	tacaggacgt	ctccccatta	caactaccca	atccgaagtg	1980
tcaaactgtg	tcaggactaa	gaacccctgg	ttttgagtag	aaaagggcct	ggaaagaggg	2040
gagccaacaa	atctgtctgc	ttcctcacat	tagtcattgg	caaataagca	ttctgtctct	2100
ttggctgctg	cctcagcaca	gagagccaga	actctatcgg	gcaccaggat	aacatctctc	2160
agtgaacaga	gttgacaagg	cctatgggaa	atgcctgatg	ggattatctt	cagcttggtg	2220
agcttctaag	tttctttccc	ttcattctac	cctgcaagcc	aagttctgta	agagaaatgc	2280
ctgagttcta	gctcagggtt	tcttactctg	aatttagatc	tccagaccct	tcctggccac	2340
aattcaaatt	aaggcaacaa	acatatacct	tccatgaagc	acacacagac	ttttgaaagc	2400
aaggacaatg	actgcttgaa	ttgaggcctt	gaggaatgaa	gctttgaagg	aaaagaatac	2460
tttgtttcca	gcccccttcc	cacactcttc	atgtgttaac	cactgccttc	ctggaccttg	2520
gagccacggg	gactgtatta	catgttggtt	tagaaaactg	attttagagt	tctgatcggt	2580
caagagaatg	attaaatata	catttcctaa	aaaaaaaaaa	aaaaaa		2626

<210> 10

<211> 1675

<212> DNA

<213> Homo sapiens

<220>

<221> SITE

<222> (1549)

<223> n equals a,t,g, or c

<400> 10

gtacgacyca	ctatagggwg	agagctatga	cgctcgcatgc	acgcgtaasc	ttgggcccct	60
cgagggatcc	tctagagcgg	ccgccctttt	tttttttttt	tttgaagaat	aacaattgag	120
ttttattctt	taaaggcatt	ctctgattta	catgagaatt	gagaaactga	gatgtatgat	180
ttgtctgtta	gtcaatttca	caccctttca	ttctcataag	cccaaattt	tgctcagtta	240
aggagcttgc	tttagggcca	cctatgtaag	tctgttatac	tagctaattg	gcccatttga	300
atagttcaag	ggtcagctaa	tgctctgagc	ttcatggctc	cagtataaag	aacaaattta	360
acaaaattaa	gctgttactg	tagccgagtt	acccttctgc	tccacacata	tgtagtggga	420
tcttgcagga	tttccatagt	gccaattatc	aaaggccttg	actacttagc	attgctgtat	480
tacagatgtg	caaactgagg	cactgaaaag	tcaaatttaa	agtcataattg	agggccagaa	540
aaggaggctt	agtttggggc	tttggccatt	ttagctactt	atctgaaatt	gctgcagata	600
caacgtatga	gcatatcaaa	tatttttgac	tgatatataat	tgattttctaa	ggtaaaaaaca	660
aataaaaaaga	aaccaataat	ttttaaaagg	aagatgtagt	tcaaaaaaaa	aaccaccatt	720
aaacatgggtg	ccattacagg	ttaaaacaaa	tgctttgtga	cttagacctc	aaaaacagag	780
cttgatgact	ttactccaca	atttgtgcac	ttagtgtata	tttaaatgct	ctctgttaat	840
tagaacaact	tcattatgct	atcaagattc	cagtaaatcca	taaaacatgt	caattatgat	900
ttgagtttgt	gcgaagccct	gtctgtgagc	tcatagtctc	aatagcctct	tctagtaccc	960

agaggaagct	atagataaaa	aataactcta	ttggcaaccc	atctgtttct	gttactggaa	1020
atttccacac	acctctgctt	ttggaaatca	cttagaaaac	ttgaggggaa	ataattcctt	1080
ttgctttcag	tctggcagca	agaaggatcc	tgaaggaatt	ctgtgggtcc	aggatccagt	1140
ggggtaattc	tgtaaagtgc	agtagtgctt	gcttaaagcc	ataggctcca	gaggtgagtc	1200
cagatcagtg	aaggggcaag	tttcatggcc	aggtgttggc	tagtcttggt	gcaggtttca	1260
gattaaagtg	ctgggtcatc	caaaggcatt	tgaaaagtgc	aaatggcaag	ctctgcaggc	1320
caccgaattc	ttgttcagag	tccagaagct	tcttttagatg	tcatatcagg	tcaccctggc	1380
tcccaagacc	acaggttcag	atagcactgt	tcacttcctt	ctttgttggtg	gtgacaggtc	1440
tttttgttgt	gtcttttgaa	gaatacagct	tttgacagag	ttgttttctt	agggcrttca	1500
ckgkggctat	gaaaatgaaa	gcaatgatgc	aggaggggat	gaaaatgtna	agcagccaag	1560
ttggatgggt	cctgggttcc	atctgacttt	gaaggtcaat	gctggccaaa	gtaagttccc	1620
tcacgtgagt	attccagaac	acacagctga	agtttctgcc	aggggttggc	tttag	1675

<210> 11

<211> 786

<212> DNA

<213> Homo sapiens

<220>

<221> SITE

<222> (754)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (778)

<223> n equals a,t,g, or c

<400> 11

ggaatgaaca	acttttcttc	tcttgaatat	atcttaacgc	caaattttga	gtgctttttt	60
gttaccatc	ctcatatgtc	ccagctggaa	agaatcctgg	gttgagacta	ctgcatgttg	120
attgttttgt	ttttcctttt	ggctgttcat	tttgggtggc	actataagga	aatctaacac	180
aaacagcaac	tgttttttgt	tgtttacttt	tgcatcttta	cttgtggagc	tgtggcaagt	240
cctcatatca	aatacagaac	atgatcttcc	tcctgcta	gttgagcctg	gaattgcagc	300
ttcaccagat	agcagcttta	ttcacagtga	cagtccttaa	ggaactgtac	ataatagagc	360
atggcagcaa	tgtgaccctg	gaatgcaact	ttgacactgg	aagtcagtgtg	aaccttggag	420
caataacagc	cagtttgcaa	aaggtggaaa	atgatacatc	cccacaccgt	gaaagagcca	480
ctttgctgga	ggagcagctg	cccctaggga	aggcctcggt	cccatmctc	aagtycaagt	540
gagggacgaa	ggacagtacc	aatgcataat	catctatggg	gtcgcctggg	actacaagta	600
cctgactctg	aaagtcaaag	cttcctacag	gaaaataaac	actcacatcc	taaaggttcc	660
agaacacagat	gaggtagagc	tcacctgcca	ggctacaggt	tatcctctgg	cagaagtatc	720
ctggccaaac	gtcagcggtc	ctgccaacac	cagncactcc	aggaccctg	aaggcctnta	780
ccaggt						786

<210> 12

<211> 2008

<212> DNA

<213> Homo sapiens

<400> 12

cgggggcttt	ctaacgggaa	aaactctact	aaagggttca	aaagctggag	ctccaccgag	60
gtggcgccg	ctctagaact	agtggatccc	ccgggctgca	ggaattcggc	acgagctcgt	120
gccgaattcg	gcacgagtca	cagaacacat	ccatggctct	matgctcagt	ttggttctga	180
gtctcctcaa	gctgggwtca	gggcagtggc	aggtgttttg	gccagacaag	cctgtccagg	240
ccttgggtgg	ggaggacgca	gcattctcct	gtttcctgtc	tcctaagacc	aatgcagagg	300
ccatggaagt	gcggttcttc	aggggccagt	tctctagcgt	ggtccacctc	tacagggacg	360
ggaaggacca	gccatttatg	cagatgccac	agtatcaagg	caggacaaaa	ctggtgaagg	420
attctattgc	ggaggggcgc	atctctctga	ggctggaaaa	cattactgtg	ttggatgctg	480
gcctctatgg	gtgcaggatt	agttcccagt	cttactacca	gaaggccatc	tgggagctac	540

agggtgtcagc	actggggtca	gttcctctca	tttccatcac	gggatatggt	gatagagaca	600
tccagctact	ctgtcagtc	tcgggctgg	tccccggcc	cacagcgaag	tggaaaggtc	660
cacaaggaca	ggatttgtcc	acagactcca	ggacaaacag	agacatgcat	ggcctgtttg	720
atgtggagat	ctctctgacc	gtccaagaga	acgccgggag	catatcctgt	tccatgcggc	780
atgtcatct	gagccgagag	gtggaatcca	gggtacagat	aggagatacc	tttttcgagc	840
ctatatcgtg	gmacttgyt	accaaagtac	tgggaatact	ctgtctggc	ctattttttg	900
gcattgttgg	actgaagatt	ttcttctcca	aattccagtg	gaaaatccag	gcggaactgg	960
actggagaag	aaagcacgga	caggcagaat	tgagagacgc	ccggaaacac	gcagtggagg	1020
tgactctgga	tccagagacg	gctcaccgga	agctctgcgt	ttctgatctg	aaaactgtaa	1080
cccatagaaa	agctccccag	gaggtgcctc	actctgagaa	gagatttaca	aggaagagtg	1140
tgggtggcttc	tcagagtttc	caagcaggga	aacattactg	ggaggtggac	ggaggacaca	1200
ataaaagggtg	gcgcgtggga	gtgtgccggg	atgatgtgga	caggaggaag	gagtacgtga	1260
ctttgtctcc	cgatcatggg	tactgggtcc	tcagactgaa	tggagaacat	ttgtatttca	1320
cattaaatcc	ccgttttatc	agcgtcttcc	ccaggacccc	acctacaaaa	ataggggtct	1380
tcctggacta	tgagtgtggg	accatctcct	tcttcaacat	aaatgaccag	tcccttattt	1440
ataccctgac	atgtcggttt	gaaggcttat	tgaggcccta	cattgagtat	ccgtcctata	1500
atgagcaaaa	tggaaactccc	agagacaagc	aacagttagt	cctcctcaca	ggcaaccacg	1560
cccttcctcc	ccaggggtga	aatgtaggat	gaatcacatc	ccacattctt	ctttagggat	1620
attaagggtct	ctctcccaga	tccaaagtcc	cgcagcagcc	ggccaagggtg	gcttccagat	1680
gaagggggac	tggcctgtcc	acatgggagt	aggtgtcat	ggctgccctg	agctgggagg	1740
gaagaaggct	gacattacat	ttagtgtgct	ctcactccat	ctggctaagt	gatcttgaaa	1800
taccacctct	cagggtgaaga	accgtcagga	attcccactc	cacaggctgt	ggtgtagatt	1860
aagtagacaa	ggaatgtgaa	taatgcttag	atcttattga	tgacagagtg	tatcctaattg	1920
gtttgttcat	tatattacac	tttcagtaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaamc	1980
tcgagggggg	gccccgtacc	caattcgg				2008

<210> 13

<211> 2799

<212> DNA

<213> Homo sapiens

<400> 13

tgggaactctg	tggaaagccca	gagaatctga	tccccgggtcc	cacaacttca	catatcgcca	60
gtaagtggga	ggcaaagaaa	attctttttc	tcctcttttg	ggacagtttg	tgactagtaa	120
tgctgtgcc	cctggaaagg	ttggagactt	gggggacgac	tggagaattg	ccatttgagg	180
accaaaggag	aaaagaaact	acacgcta	tctagaaggc	ctcctgtccc	tgctgtctct	240
gggtgtcat	ggaaccagct	gctgccctgc	acttctcccg	gccagcctcc	ctcctcctcc	300
tcctcagcct	gtgtgcactg	gtctcagccc	agtttactgt	cgtagggcca	gctaattcca	360
tcctggccat	ggtgggagaa	aacactacat	tacgtgcgca	tctgtcacc	gagaaaaatg	420
ctgaggacat	ggaggtgcgg	tggttccggg	ctcagttctc	ccccgcagtg	tttgtgtata	480
aggggtgggag	agagagaaca	gaggagcaga	tggaggagta	ccggggaaga	atcacctttg	540
tgagcaaaaga	catcaacagg	ggcagcgtgg	ccctggctcat	acataacgtc	acagcccagg	600
agaatgggat	ctaccgctgt	tacttccaag	aaggcaggtc	ctacgatgag	gccatcctac	660
gcctcgtggg	ggcaggcctt	gggtctaagc	ccctcattga	aatcaaggcc	caagaggatg	720
ggagcatctg	gctggagtgc	atatctggag	ggtggtaccc	agagcccctc	acagtgtgga	780
gggaccccta	cggtgagggt	gtgcccgccc	tgaaggagggt	ttccatcgct	gatgtgcacg	840
gcctcttcat	ggtcaccaca	gctgtgatca	tcagagacaa	gtatgtgagg	aatgtgtcct	900
gctctgtcaa	caacaccctg	ctcgccagg	agaaggaaac	tgtcattttt	attccagaat	960
ccttttatgcc	cagcgcactc	ccctggatgg	tggccctagc	tgtcatcctg	accgcattctc	1020
cctggatggg	gtccatgact	gtcatcctgg	ctgttttcat	catcttcatg	gctgtcagca	1080
tctgttgc	caagaaactt	caaaggga	aaaagattct	gtcaggggaa	aagaaagtgtg	1140
aacaagagga	aaaagaaatt	gcacagcaac	ttcaagaaga	attgcgatgg	agaagaacat	1200
tcttacatgc	tgctgatgtg	gtcctggatc	cagacaccgc	tcatcccag	ctcttctgt	1260
cagaggaccg	gagaagtgtg	aggcggggcc	cctacaggca	gagagtgcct	gacaaccacg	1320
agagattcga	cagtcagcct	tgtgtcctgg	gatgggagag	cttcgcctca	gggaaacatt	1380
acagggga	cttcacagag	tggggaccct	ccagagccta	tagaatcaat	tccttggact	1440
cacagccatg	cagataagcc	ctggccatct	cagcagccac	cgcaacaacc	ccctaataa	1500
agacacgccc	tcctcccctc	tggtcacgta	agagacatc	ttccagctgc	ctttttcaca	1560
ccactccag	ccctctgccc	cagttttctc	ctctcacta	gtctgtgggt	ttagtgttc	1620
ctttgttgt	aattatggga	tgggatccag	gcatagggaa	ctagtgtgtt	catagctccc	1680

```

agtcaaaaaag aaagtgagag aagctgttg gcagcgaacc tactgtttaa aatcaggata 1740
accacattaa gcccaatat ccagttggca ccagatgctg tggacttga atgaggccaa 1800
cagggttcac caggatgaga gaggagagag gaatccacag gaccaccaga agggagaggg 1860
aaccagatat gcagatcaga gatagaggaa gtgttgagag gaaaggggag gtcctgctga 1920
ttcctcagaa tggcttctgg accctggaga tgtttggaaa ccaataccgg gccctgtcct 1980
cccctgagag gattctccct ttgaaggagt ccctttgcg ggtgggcgtc ttcctggact 2040
atgaagctgg agatgtctcc ttctacaaca tgagggacag atcacacatc tacacatgctc 2100
cccgttcagc ctttactgtg cctgtgaggc ccttcttcag gttagggtct gatgacagcc 2160
ccatcttcat ctgccctgca ctcacaggag ccagtggggg catggtgcct gaagagggcc 2220
tgaaacttca cagagtgggg acccaccaag gttgtaaggg atggctaagt cccaccataa 2280
gagctaaagg gtcctgggag atgatggctc atttccaccc aaccccagga tttccacagc 2340
acacaccac aggcctggac ctgggatgaa gatgaatgaa gaacatggac tcatgtggat 2400
gtgggtttggc tcagatgtcc ctgcaataaa caaggggtca gtacttagtc cctgagtgtg 2460
gttgaggttt gaggtcctgg tcgagcaggg cagtactgga ccaggtctac gtcagcattc 2520
aggttcaatg ggggacacca gtggcttcaa acttctgat ctaattatgt ttttagacac 2580
ttagaagtta ttgaggactt taaagaactt ttgtttattt gggttaatat ttatgacatt 2640
tgaccattga aacaaaaatt taaaatgtta tcttttaatt tatgttaaaa tagcattaat 2700
aatcagtta taggttaatg tagataggat gttttgtgaa aaagcaatct attgtgtcca 2760
aataaaaaaa caaaaagtgt aaaaaaaaaa aaaaaaaaaa 2799

```

<210> 14
 <211> 282
 <212> PRT
 <213> Homo sapiens

<400> 14
 Met Ala Ser Leu Gly Gln Ile Leu Phe Trp Ser Ile Ile Ser Ile Ile
 1 5 10 15
 Ile Ile Leu Ala Gly Ala Ile Ala Leu Ile Ile Gly Phe Gly Ile Ser
 20 25 30
 Gly Arg His Ser Ile Thr Val Thr Thr Val Ala Ser Ala Gly Asn Ile
 35 40 45
 Gly Glu Asp Gly Ile Leu Ser Cys Thr Phe Glu Pro Asp Ile Lys Leu
 50 55 60
 Ser Asp Ile Val Ile Gln Trp Leu Lys Glu Gly Val Leu Gly Leu Val
 65 70 75 80
 His Glu Phe Lys Glu Gly Lys Asp Glu Leu Ser Glu Gln Asp Glu Met
 85 90 95
 Phe Arg Gly Arg Thr Ala Val Phe Ala Asp Gln Val Ile Val Gly Asn
 100 105 110
 Ala Ser Leu Arg Leu Lys Asn Val Gln Leu Thr Asp Ala Gly Thr Tyr
 115 120 125
 Lys Cys Tyr Ile Ile Thr Ser Lys Gly Lys Gly Asn Ala Asn Leu Glu
 130 135 140
 Tyr Lys Thr Gly Ala Phe Ser Met Pro Glu Val Asn Val Asp Tyr Asn
 145 150 155 160
 Ala Ser Ser Glu Thr Leu Arg Cys Glu Ala Pro Arg Trp Phe Pro Gln
 165 170 175
 Pro Thr Val Val Trp Ala Ser Gln Val Asp Gln Gly Ala Asn Phe Ser

	180						185						190					
Glu Val Ser Asn Thr Ser Phe Glu Leu Asn Ser Glu Asn Val Thr Met 195 200 205																		
Lys Val Val Ser Val Leu Tyr Asn Val Thr Ile Asn Asn Thr Tyr Ser 210 215 220																		
Cys Met Ile Glu Asn Asp Ile Ala Lys Ala Thr Gly Asp Ile Lys Val 225 230 235 240																		
Thr Glu Ser Glu Ile Lys Arg Arg Ser His Leu Gln Leu Leu Asn Ser 245 250 255																		
Lys Ala Ser Leu Cys Val Ser Ser Phe Phe Ala Ile Ser Trp Ala Leu 260 265 270																		
Leu Pro Leu Ser Pro Tyr Leu Met Leu Lys 275 280																		

```
<210> 15
<211> 283
<212> PRT
<213> Homo sapiens
```

<400> 15																
Met	Ile	Phe	Leu	Leu	Leu	Met	Leu	Ser	Leu	Glu	Leu	Gln	Leu	His	Gln	
1				5					10					15		
Ile	Ala	Ala	Leu	Phe	Thr	Val	Thr	Val	Pro	Lys	Glu	Leu	Tyr	Ile	Ile	
			20					25					30			
Glu	His	Gly	Ser	Asn	Val	Thr	Leu	Glu	Cys	Asn	Phe	Asp	Thr	Gly	Ser	
		35					40					45				
His	Val	Asn	Leu	Gly	Ala	Ile	Thr	Ala	Ser	Leu	Gln	Lys	Val	Glu	Asn	
	50					55					60					
Asp	Thr	Ser	Pro	His	Arg	Glu	Arg	Ala	Thr	Leu	Leu	Glu	Glu	Gln	Leu	
65					70					75					80	
Pro	Leu	Gly	Lys	Ala	Ser	Phe	His	Ile	Pro	Gln	Val	Gln	Val	Arg	Asp	
				85					90					95		
Glu	Gly	Gln	Tyr	Gln	Cys	Ile	Ile	Ile	Tyr	Gly	Val	Ala	Trp	Asp	Tyr	
			100					105					110			
Lys	Tyr	Leu	Thr	Leu	Lys	Val	Lys	Ala	Ser	Tyr	Arg	Lys	Ile	Asn	Thr	
		115					120					125				
His	Ile	Leu	Lys	Val	Pro	Glu	Thr	Asp	Glu	Val	Glu	Leu	Thr	Cys	Gln	
	130					135					140					
Ala	Thr	Gly	Tyr	Pro	Leu	Ala	Glu	Val	Ser	Trp	Pro	Asn	Val	Ser	Val	
145					150					155					160	
Pro	Ala	Asn	Thr	Ser	His	Ser	Arg	Thr	Pro	Glu	Gly	Leu	Tyr	Gln	Val	
				165					170					175		
Thr	Ser	Val	Leu	Arg	Leu	Lys	Pro	Pro	Pro	Gly	Arg	Asn	Phe	Ser	Cys	

180 185 190
 Val Phe Trp Asn Thr His Val Arg Glu Leu Thr Leu Ala Ser Ile Asp
 195 200 205
 Leu Gln Ser Gln Met Glu Pro Arg Thr His Pro Thr Trp Leu Leu His
 210 215 220
 Ile Phe Ile Pro Ser Cys Ile Ile Ala Phe Ile Phe Ile Ala Thr Val
 225 230 235 240
 Ile Ala Leu Arg Lys Gln Leu Cys Gln Lys Leu Tyr Ser Ser Lys Asp
 245 250 255
 Thr Thr Lys Arg Pro Val Thr Thr Thr Lys Arg Glu Val Asn Ser Ala
 260 265 270
 Val Asn Leu Asn Leu Trp Ser Trp Glu Pro Gly
 275 280

<210> 16
 <211> 318
 <212> PRT
 <213> Homo sapiens

<400> 16
 Met Ala Leu Met Leu Ser Leu Val Leu Ser Leu Leu Lys Leu Gly Ser
 1 5 10 15
 Gly Gln Trp Gln Val Phe Gly Pro Asp Lys Pro Val Gln Ala Leu Val
 20 25 30
 Gly Glu Asp Ala Ala Phe Ser Cys Phe Leu Ser Pro Lys Thr Asn Ala
 35 40 45
 Glu Ala Met Glu Val Arg Phe Phe Arg Gly Gln Phe Ser Ser Val Val
 50 55 60
 His Leu Tyr Arg Asp Gly Lys Asp Gln Pro Phe Met Gln Met Pro Gln
 65 70 75 80
 Tyr Gln Gly Arg Thr Lys Leu Val Lys Asp Ser Ile Ala Glu Gly Arg
 85 90 95
 Ile Ser Leu Arg Leu Glu Asn Ile Thr Val Leu Asp Ala Gly Leu Tyr
 100 105 110
 Gly Cys Arg Ile Ser Ser Gln Ser Tyr Tyr Gln Lys Ala Ile Trp Glu
 115 120 125
 Leu Gln Val Ser Ala Leu Gly Ser Val Pro Leu Ile Ser Ile Ala Gly
 130 135 140
 Tyr Val Asp Arg Asp Ile Gln Leu Leu Cys Gln Ser Ser Gly Trp Phe
 145 150 155 160
 Pro Arg Pro Thr Ala Lys Trp Lys Gly Pro Gln Gly Gln Asp Leu Ser
 165 170 175
 Thr Asp Ser Arg Thr Asn Arg Asp Met His Gly Leu Phe Asp Val Glu

180 185 190
 Ile Ser Leu Thr Val Gln Glu Asn Ala Gly Ser Ile Ser Cys Ser Met
 195 200 205
 Arg His Ala His Leu Ser Arg Glu Val Glu Ser Arg Val Gln Ile Gly
 210 215 220
 Asp Trp Arg Arg Lys His Gly Gln Ala Gly Lys Arg Lys Tyr Ser Ser
 225 230 235 240
 Ser His Ile Tyr Asp Ser Phe Pro Ser Leu Ser Phe Met Asp Phe Tyr
 245 250 255
 Ile Leu Arg Pro Val Gly Pro Cys Arg Ala Lys Leu Val Met Gly Thr
 260 265 270
 Leu Lys Leu Gln Ile Leu Gly Glu Val His Phe Val Glu Lys Pro His
 275 280 285
 Ser Leu Leu Gln Ile Ser Gly Gly Ser Thr Thr Leu Lys Lys Gly Pro
 290 295 300
 Asn Pro Trp Ser Phe Pro Ser Pro Cys Ala Leu Phe Pro Thr
 305 310 315

<210> 17
 <211> 454
 <212> PRT
 <213> Homo sapiens

<400> 17
 Met Glu Pro Ala Ala Ala Leu His Phe Ser Arg Pro Ala Ser Leu Leu
 1 5 10 15
 Leu Leu Leu Ser Leu Cys Ala Leu Val Ser Ala Gln Phe Thr Val Val
 20 25 30
 Gly Pro Ala Asn Pro Ile Leu Ala Met Val Gly Glu Asn Thr Thr Leu
 35 40 45
 Arg Cys His Leu Ser Pro Glu Lys Asn Ala Glu Asp Met Glu Val Arg
 50 55 60
 Trp Phe Arg Ser Gln Phe Ser Pro Ala Val Phe Val Tyr Lys Gly Gly
 65 70 75 80
 Arg Glu Arg Thr Glu Glu Gln Met Glu Glu Tyr Arg Gly Arg Ile Thr
 85 90 95
 Phe Val Ser Lys Asp Ile Asn Arg Gly Ser Val Ala Leu Val Ile His
 100 105 110
 Asn Val Thr Ala Gln Glu Asn Gly Ile Tyr Arg Cys Tyr Phe Gln Glu
 115 120 125
 Gly Arg Ser Tyr Asp Glu Ala Ile Leu Arg Leu Val Val Ala Gly Leu
 130 135 140
 Gly Ser Lys Pro Leu Ile Glu Ile Lys Ala Gln Glu Asp Gly Ser Ile

145		150		155		160									
Trp	Leu	Glu	Cys	Ile	Ser	Gly	Gly	Trp	Tyr	Pro	Glu	Pro	Leu	Thr	Val
				165					170					175	
Trp	Arg	Asp	Pro	Tyr	Gly	Glu	Val	Val	Pro	Ala	Leu	Lys	Glu	Val	Ser
			180					185					190		
Ile	Ala	Asp	Ala	Asp	Gly	Leu	Phe	Met	Val	Thr	Thr	Ala	Val	Ile	Ile
		195					200					205			
Arg	Asp	Lys	Tyr	Val	Arg	Asn	Val	Ser	Cys	Ser	Val	Asn	Asn	Thr	Leu
	210					215					220				
Leu	Gly	Gln	Glu	Lys	Glu	Thr	Val	Ile	Phe	Ile	Pro	Glu	Ser	Phe	Met
225					230				235						240
Pro	Ser	Ala	Ser	Pro	Trp	Met	Val	Ala	Leu	Ala	Val	Ile	Leu	Thr	Ala
				245					250					255	
Ser	Pro	Trp	Met	Val	Ser	Met	Thr	Val	Ile	Leu	Ala	Val	Phe	Ile	Ile
			260					265					270		
Phe	Met	Ala	Val	Ser	Ile	Cys	Cys	Ile	Lys	Lys	Leu	Gln	Arg	Glu	Lys
	275						280					285			
Lys	Ile	Leu	Ser	Gly	Glu	Lys	Lys	Val	Glu	Gln	Glu	Glu	Lys	Glu	Ile
	290					295					300				
Ala	Gln	Gln	Leu	Gln	Glu	Glu	Leu	Arg	Trp	Arg	Arg	Thr	Phe	Leu	His
305					310					315					320
Ala	Ala	Asp	Val	Val	Leu	Asp	Pro	Asp	Thr	Ala	His	Pro	Glu	Leu	Phe
				325					330					335	
Leu	Ser	Glu	Asp	Arg	Arg	Ser	Val	Arg	Arg	Gly	Pro	Tyr	Arg	Gln	Arg
			340					345					350		
Val	Pro	Asp	Asn	Pro	Glu	Arg	Phe	Asp	Ser	Gln	Pro	Cys	Val	Leu	Gly
		355					360					365			
Trp	Glu	Ser	Phe	Ala	Ser	Gly	Lys	His	Tyr	Arg	Gly	Asn	Phe	Thr	Glu
	370					375					380				
Trp	Gly	Pro	Thr	Arg	Ala	Tyr	Arg	Ile	Asn	Ser	Leu	Asp	Ser	Gln	Pro
385					390					395					400
Cys	Arg	Lys	Pro	Trp	Pro	Ser	Gln	Gln	Pro	Pro	His	Asn	Pro	Pro	Asn
				405					410					415	
Glu	Arg	His	Ala	Leu	Leu	Pro	Ser	Gly	His	Val	Arg	Glu	His	Leu	Pro
			420					425					430		
Ala	Ala	Phe	Phe	Thr	Pro	Thr	Pro	Ala	Leu	Cys	Pro	Ser	Phe	Leu	Leu
		435					440					445			
Leu	Thr	Ser	Leu	Trp	Leu										
	450														

<210> 18

<211> 414

<212> PRT

<213> Homo sapiens

<400> 18

```

Met Arg Glu Ile Val Trp Tyr Arg Val Thr Asp Gly Gly Thr Ile Lys
 1           5           10           15

Gln Lys Ile Phe Thr Phe Asp Ala Met Phe Ser Thr Asn Tyr Ser His
          20           25           30

Met Glu Asn Tyr Arg Lys Arg Glu Asp Leu Val Tyr Gln Ser Thr Val
 35           40           45

Arg Leu Pro Glu Val Arg Ile Ser Asp Asn Gly Pro Tyr Glu Cys His
 50           55           60

Val Gly Ile Tyr Asp Arg Ala Thr Arg Glu Lys Val Val Leu Ala Ser
 65           70           75           80

Gly Asn Ile Phe Leu Asn Val Met Ala Pro Pro Thr Ser Ile Glu Val
          85           90           95

Val Ala Ala Asp Thr Pro Ala Pro Phe Ser Arg Tyr Gln Ala Gln Asn
          100          105          110

Phe Thr Leu Val Cys Ile Val Ser Gly Gly Lys Pro Ala Pro Met Val
          115          120          125

Tyr Phe Lys Arg Asp Gly Glu Pro Ile Asp Ala Val Pro Leu Ser Glu
          130          135          140

Pro Pro Ala Ala Ser Ser Gly Pro Leu Gln Asp Ser Arg Pro Phe Arg
          145          150          155          160

Ser Leu Leu His Arg Asp Leu Asp Asp Thr Lys Met Gln Lys Ser Leu
          165          170          175

Ser Leu Leu Asp Ala Glu Asn Arg Gly Gly Arg Pro Tyr Thr Glu Arg
          180          185          190

Pro Ser Arg Gly Leu Thr Pro Asp Pro Asn Ile Leu Leu Gln Pro Thr
          195          200          205

Thr Glu Asn Ile Pro Glu Thr Val Val Ser Arg Glu Phe Pro Arg Trp
          210          215          220

Val His Ser Ala Glu Pro Thr Tyr Phe Leu Arg His Ser Arg Thr Pro
          225          230          235          240

Ser Ser Asp Gly Thr Val Glu Val Arg Ala Leu Leu Thr Trp Thr Leu
          245          250          255

Asn Pro Gln Ile Asp Asn Glu Ala Leu Phe Ser Cys Glu Val Lys His
          260          265          270

Pro Ala Leu Ser Met Pro Met Gln Ala Glu Val Thr Leu Val Ala Pro
          275          280          285

Lys Gly Pro Lys Ile Val Met Thr Pro Ser Arg Ala Arg Val Gly Asp
          290          295          300

```

Thr Val Arg Ile Leu Val His Gly Phe Gln Asn Glu Val Phe Pro Glu
 305 310 315 320
 Pro Met Phe Thr Trp Thr Arg Val Gly Ser Arg Leu Leu Asp Gly Ser
 325 330 335
 Ala Glu Phe Asp Gly Lys Glu Leu Val Leu Glu Arg Val Pro Ala Glu
 340 345 350
 Leu Asn Gly Ser Met Tyr Arg Cys Thr Ala Gln Asn Pro Leu Gly Ser
 355 360 365
 Thr Asp Thr His Thr Arg Leu Ile Val Phe Glu Asn Pro Asn Ile Pro
 370 375 380
 Arg Gly Thr Glu Asp Ser Asn Gly Ser Ile Gly Pro Thr Gly Ala Arg
 385 390 395 400
 Leu Thr Leu Val Leu Ala Leu Thr Val Ile Leu Glu Leu Thr
 405 410

<210> 19
 <211> 159
 <212> PRT
 <213> Homo sapiens

<400> 19
 Met Glu Pro Ala Ala Ala Leu His Phe Ser Arg Pro Ala Ser Leu Leu
 1 5 10 15
 Leu Leu Leu Ser Leu Cys Ala Leu Val Ser Ala Gln Val Thr Val Val
 20 25 30
 Gly Pro Thr Asp Pro Ile Leu Ala Met Val Gly Glu Asn Thr Thr Leu
 35 40 45
 Arg Cys Cys Leu Ser Pro Glu Glu Asn Ala Glu Asp Met Glu Val Arg
 50 55 60
 Trp Phe Gln Ser Gln Phe Ser Pro Ala Val Phe Val Tyr Lys Gly Gly
 65 70 75 80
 Arg Glu Arg Thr Glu Glu Gln Lys Glu Glu Tyr Arg Gly Arg Thr Thr
 85 90 95
 Phe Val Ser Lys Asp Ser Arg Gly Ser Val Ala Leu Ile Ile His Asn
 100 105 110
 Val Thr Ala Glu Asp Asn Gly Ile Tyr Gln Cys Tyr Phe Gln Glu Gly
 115 120 125
 Arg Ser Cys Asn Glu Ala Ile Leu His Leu Val Val Ala Asp Gln His
 130 135 140
 Asn Pro Leu Ser Trp Ile Pro Ile Pro Gln Gly Thr Leu Ser Leu
 145 150 155

<210> 20

<211> 461

<212> PRT

<213> Homo sapiens

<400> 20

Met Ala Leu Met Leu Ser Leu Val Leu Ser Leu Leu Lys Leu Gly Ser
 1 5 10 15

Gly Gln Trp Gln Val Phe Gly Pro Asp Lys Pro Val Gln Ala Leu Val
 20 25 30

Gly Glu Asp Ala Ala Phe Ser Cys Phe Leu Ser Pro Lys Thr Asn Ala
 35 40 45

Glu Ala Met Glu Val Arg Phe Phe Arg Gly Gln Phe Ser Ser Val Val
 50 55 60

His Leu Tyr Arg Asp Gly Lys Asp Gln Pro Phe Met Gln Met Pro Gln
 65 70 75 80

Tyr Gln Gly Arg Thr Lys Leu Val Lys Asp Ser Ile Ala Glu Gly Arg
 85 90 95

Ile Ser Leu Arg Leu Glu Asn Ile Thr Val Leu Asp Ala Gly Leu Tyr
 100 105 110

Gly Cys Arg Ile Ser Ser Gln Ser Tyr Tyr Gln Lys Ala Ile Trp Glu
 115 120 125

Leu Gln Val Ser Ala Leu Gly Ser Val Pro Leu Ile Ser Ile Thr Gly
 130 135 140

Tyr Val Asp Arg Asp Ile Gln Leu Leu Cys Gln Ser Ser Gly Trp Phe
 145 150 155 160

Pro Arg Pro Thr Ala Lys Trp Lys Gly Pro Gln Gly Gln Asp Leu Ser
 165 170 175

Thr Asp Ser Arg Thr Asn Arg Asp Met His Gly Leu Phe Asp Val Glu
 180 185 190

Ile Ser Leu Thr Val Gln Glu Asn Ala Gly Ser Ile Ser Cys Ser Met
 195 200 205

Arg His Ala His Leu Ser Arg Glu Val Glu Ser Arg Val Gln Ile Gly
 210 215 220

Asp Thr Phe Phe Glu Pro Ile Ser Trp His Leu Ala Thr Lys Val Leu
 225 230 235 240

Gly Ile Leu Cys Cys Gly Leu Phe Phe Gly Ile Val Gly Leu Lys Ile
 245 250 255

Phe Phe Ser Lys Phe Gln Trp Lys Ile Gln Ala Glu Leu Asp Trp Arg
 260 265 270

Arg Lys His Gly Gln Ala Glu Leu Arg Asp Ala Arg Lys His Ala Val
 275 280 285

Glu Val Thr Leu Asp Pro Glu Thr Ala His Pro Lys Leu Cys Val Ser
 290 295 300

Asp Leu Lys Thr Val Thr His Arg Lys Ala Pro Gln Glu Val Pro His
305 310 315 320

Ser Glu Lys Arg Phe Thr Arg Lys Ser Val Val Ala Ser Gln Ser Phe
325 330 335

Gln Ala Gly Lys His Tyr Trp Glu Val Asp Gly Gly His Asn Lys Arg
340 345 350

Trp Arg Val Gly Val Cys Arg Asp Asp Val Asp Arg Arg Lys Glu Tyr
355 360 365

Val Thr Leu Ser Pro Asp His Gly Tyr Trp Val Leu Arg Leu Asn Gly
370 375 380

Glu His Leu Tyr Phe Thr Leu Asn Pro Arg Phe Ile Ser Val Phe Pro
385 390 395 400

Arg Thr Pro Pro Thr Lys Ile Gly Val Phe Leu Asp Tyr Glu Cys Gly
405 410 415

Thr Ile Ser Phe Phe Asn Ile Asn Asp Gln Ser Leu Ile Tyr Thr Leu
420 425 430

Thr Cys Arg Phe Glu Gly Leu Leu Arg Pro Tyr Ile Glu Tyr Pro Ser
435 440 445

Tyr Asn Glu Gln Asn Gly Thr Pro Arg Asp Lys Gln Gln
450 455 460

<210> 21

<211> 13

<212> PRT

<213> Homo sapiens

<400> 21

Met Ala Ser Leu Gly Gln Ile Leu Phe Trp Ser Ile Ile
1 5 10

<210> 22

<211> 23

<212> PRT

<213> Homo sapiens

<400> 22

Leu Phe Leu Leu Leu Glu Ile Ser Thr His Leu Cys Phe Trp Lys Ser
1 5 10 15

Leu Arg Lys Leu Glu Gly Lys
20

<210> 23

<211> 93

<212> PRT

<213> Homo sapiens

<220>

<221> SITE

<222> (89)

<223> Xaa equals any of the naturally occurring L-amino acids

<220>

<221> SITE

<222> (92)

<223> Xaa equals any of the naturally occurring L-amino acids

<400> 23

Met	Ile	Phe	Leu	Leu	Leu	Met	Leu	Ser	Leu	Glu	Leu	Gln	Leu	His	Gln
1				5					10					15	

Ile	Ala	Ala	Leu	Phe	Thr	Val	Thr	Val	Pro	Lys	Glu	Leu	Tyr	Ile	Ile
			20					25					30		

Glu	His	Gly	Ser	Asn	Val	Thr	Leu	Glu	Cys	Asn	Phe	Asp	Thr	Gly	Ser
		35					40					45			

His	Val	Asn	Leu	Gly	Ala	Ile	Thr	Ala	Ser	Leu	Gln	Lys	Val	Glu	Asn
	50					55					60				

Asp	Thr	Ser	Pro	His	Arg	Glu	Arg	Ala	Thr	Leu	Leu	Glu	Glu	Gln	Leu
65					70					75					80

Pro	Leu	Gly	Lys	Ala	Ser	Phe	Pro	Xaa	Leu	Lys	Xaa	Lys
			85						90			

<210> 24

<211> 461

<212> PRT

<213> Homo sapiens

<220>

<221> SITE

<222> (234)

<223> Xaa equals any of the naturally occurring L-amino acids

<220>

<221> SITE

<222> (236)

<223> Xaa equals any of the naturally occurring L-amino acids

<400> 24

Met	Ala	Leu	Met	Leu	Ser	Leu	Val	Leu	Ser	Leu	Leu	Lys	Leu	Gly	Ser
1				5					10					15	

Gly	Gln	Trp	Gln	Val	Phe	Gly	Pro	Asp	Lys	Pro	Val	Gln	Ala	Leu	Val
			20					25					30		

Gly	Glu	Asp	Ala	Ala	Phe	Ser	Cys	Phe	Leu	Ser	Pro	Lys	Thr	Asn	Ala
		35					40					45			

Glu	Ala	Met	Glu	Val	Arg	Phe	Phe	Arg	Gly	Gln	Phe	Ser	Ser	Val	Val
	50					55				60					

His	Leu	Tyr	Arg	Asp	Gly	Lys	Asp	Gln	Pro	Phe	Met	Gln	Met	Pro	Gln
65					70					75					80

Tyr	Gln	Gly	Arg	Thr	Lys	Leu	Val	Lys	Asp	Ser	Ile	Ala	Glu	Gly	Arg
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

21

Thr Ile Ser Phe Phe Asn Ile Asn Asp Gln Ser Leu Ile Tyr Thr Leu
 420 425 430

Thr Cys Arg Phe Glu Gly Leu Leu Arg Pro Tyr Ile Glu Tyr Pro Ser
 435 440 445

Tyr Asn Glu Gln Asn Gly Thr Pro Arg Asp Lys Gln Gln
 450 455 460

<210> 25
 <211> 402
 <212> PRT
 <213> Homo sapiens

<400> 25
 Met Glu Pro Ala Ala Ala Leu His Phe Ser Arg Pro Ala Ser Leu Leu
 1 5 10 15

Leu Leu Leu Ser Leu Cys Ala Leu Val Ser Ala Gln Phe Thr Val Val
 20 25 30

Gly Pro Ala Asn Pro Ile Leu Ala Met Val Gly Glu Asn Thr Thr Leu
 35 40 45

Arg Cys His Leu Ser Pro Glu Lys Asn Ala Glu Asp Met Glu Val Arg
 50 55 60

Trp Phe Arg Ser Gln Phe Ser Pro Ala Val Phe Val Tyr Lys Gly Gly
 65 70 75 80

Arg Glu Arg Thr Glu Glu Gln Met Glu Glu Tyr Arg Gly Arg Ile Thr
 85 90 95

Phe Val Ser Lys Asp Ile Asn Arg Gly Ser Val Ala Leu Val Ile His
 100 105 110

Asn Val Thr Ala Gln Glu Asn Gly Ile Tyr Arg Cys Tyr Phe Gln Glu
 115 120 125

Gly Arg Ser Tyr Asp Glu Ala Ile Leu Arg Leu Val Val Ala Gly Leu
 130 135 140

Gly Ser Lys Pro Leu Ile Glu Ile Lys Ala Gln Glu Asp Gly Ser Ile
 145 150 155 160

Trp Leu Glu Cys Ile Ser Gly Gly Trp Tyr Pro Glu Pro Leu Thr Val
 165 170 175

Trp Arg Asp Pro Tyr Gly Glu Val Val Pro Ala Leu Lys Glu Val Ser
 180 185 190

Ile Ala Asp Ala Asp Gly Leu Phe Met Val Thr Thr Ala Val Ile Ile
 195 200 205

Arg Asp Lys Tyr Val Arg Asn Val Ser Cys Ser Val Asn Asn Thr Leu
 210 215 220

Leu Gly Gln Glu Lys Glu Thr Val Ile Phe Ile Pro Glu Ser Phe Met
 225 230 235 240

Pro	Ser	Ala	Ser	Pro	Trp	Met	Val	Ala	Leu	Ala	Val	Ile	Leu	Thr	Ala	
				245					250							255
Ser	Pro	Trp	Met	Val	Ser	Met	Thr	Val	Ile	Leu	Ala	Val	Phe	Ile	Ile	
				260					265							270
Phe	Met	Ala	Val	Ser	Ile	Cys	Cys	Ile	Lys	Lys	Leu	Gln	Arg	Glu	Lys	
				275					280							285
Lys	Ile	Leu	Ser	Gly	Glu	Lys	Lys	Val	Glu	Gln	Glu	Glu	Lys	Glu	Ile	
				290					295							300
Ala	Gln	Gln	Leu	Gln	Glu	Glu	Leu	Arg	Trp	Arg	Arg	Thr	Phe	Leu	His	
				305					310							320
Ala	Ala	Asp	Val	Val	Leu	Asp	Pro	Asp	Thr	Ala	His	Pro	Glu	Leu	Phe	
				325					330							335
Leu	Ser	Glu	Asp	Arg	Arg	Ser	Val	Arg	Arg	Gly	Pro	Tyr	Arg	Gln	Arg	
				340					345							350
Val	Pro	Asp	Asn	Pro	Glu	Arg	Phe	Asp	Ser	Gln	Pro	Cys	Val	Leu	Gly	
				355					360							365
Trp	Glu	Ser	Phe	Ala	Ser	Gly	Lys	His	Tyr	Arg	Gly	Asn	Phe	Thr	Glu	
				370					375							380
Trp	Gly	Pro	Thr	Arg	Ala	Tyr	Arg	Ile	Asn	Ser	Leu	Asp	Ser	Gln	Pro	
				385					390							400
Cys		Arg														

```
<210> 26
<211> 20
<212> PRT
<213> Homo sapiens
```

```
<400> 26
Ser Lys Ala Ser Leu Cys Val Ser Ser Phe Phe Ala Ile Ser Trp Ala
  1                      5              10              15
Leu Leu Pro Leu
      20
```

```
<210> 27
<211> 255
<212> PRT
<213> Homo sapiens
```

```

<400> 27
Met Ala Ser Leu Gly Gln Ile Leu Phe Trp Ser Ile Ile Ser Ile Ile
  1             5             10             15

Ile Ile Leu Ala Gly Ala Ile Ala Leu Ile Ile Gly Phe Gly Ile Ser
      20             25             30

Gly Arg His Ser Ile Thr Val Thr Thr Val Ala Ser Ala Gly Asn Ile

```

35 40 45
 Gly Glu Asp Gly Ile Leu Ser Cys Thr Phe Glu Pro Asp Ile Lys Leu
 50 55 60
 Ser Asp Ile Val Ile Gln Trp Leu Lys Glu Gly Val Leu Gly Leu Val
 65 70 75 80
 His Glu Phe Lys Glu Gly Lys Asp Glu Leu Ser Glu Gln Asp Glu Met
 85 90 95
 Phe Arg Gly Arg Thr Ala Val Phe Ala Asp Gln Val Ile Val Gly Asn
 100 105 110
 Ala Ser Leu Arg Leu Lys Asn Val Gln Leu Thr Asp Ala Gly Thr Tyr
 115 120 125
 Lys Cys Tyr Ile Ile Thr Ser Lys Gly Lys Gly Asn Ala Asn Leu Glu
 130 135 140
 Tyr Lys Thr Gly Ala Phe Ser Met Pro Glu Val Asn Val Asp Tyr Asn
 145 150 155 160
 Ala Ser Ser Glu Thr Leu Arg Cys Glu Ala Pro Arg Trp Phe Pro Gln
 165 170 175
 Pro Thr Val Val Trp Ala Ser Gln Val Asp Gln Gly Ala Asn Phe Ser
 180 185 190
 Glu Val Ser Asn Thr Ser Phe Glu Leu Asn Ser Glu Asn Val Thr Met
 195 200 205
 Lys Val Val Ser Val Leu Tyr Asn Val Thr Ile Asn Asn Thr Tyr Ser
 210 215 220
 Cys Met Ile Glu Asn Asp Ile Ala Lys Ala Thr Gly Asp Ile Lys Val
 225 230 235 240
 Thr Glu Ser Glu Ile Lys Arg Arg Ser His Leu Gln Leu Leu Asn
 245 250 255

 <210> 28
 <211> 231
 <212> PRT
 <213> Homo sapiens

 <400> 28
 Leu Ile Ile Gly Phe Gly Ile Ser Gly Arg His Ser Ile Thr Val Thr
 1 5 10 15
 Thr Val Ala Ser Ala Gly Asn Ile Gly Glu Asp Gly Ile Leu Ser Cys
 20 25 30
 Thr Phe Glu Pro Asp Ile Lys Leu Ser Asp Ile Val Ile Gln Trp Leu
 35 40 45
 Lys Glu Gly Val Leu Gly Leu Val His Glu Phe Lys Glu Gly Lys Asp
 50 55 60
 Glu Leu Ser Glu Gln Asp Glu Met Phe Arg Gly Arg Thr Ala Val Phe

```
<210> 29
<211> 24
<212> PRT
<213> Homo sapiens
```

```

<400> 29
Met Ala Ser Leu Gly Gln Ile Leu Phe Trp Ser Ile Ile Ser Ile Ile
  1             5             10             15

Ile Ile Leu Ala Gly Ala Ile Ala
          20

```

```
<210> 30
<211> 30
<212> PRT
<213> Homo sapiens
```

```
<400> 30
Pro Thr Trp Leu Leu His Ile Phe Ile Pro Ser Cys Ile Ile Ala Phe
  1                               10                          15
Ile Phe Ile Ala Thr Val Ile Ala Leu Arg Lys Gln Leu Cys
      20                      25                     30
```

<210> 31
<211> 218

<212> PRT

<213> Homo sapiens

<400> 31

```

Met Ile Phe Leu Leu Leu Met Leu Ser Leu Glu Leu Gln Leu His Gln
 1             5             10             15

Ile Ala Ala Leu Phe Thr Val Thr Val Pro Lys Glu Leu Tyr Ile Ile
      20             25             30

Glu His Gly Ser Asn Val Thr Leu Glu Cys Asn Phe Asp Thr Gly Ser
      35             40             45

His Val Asn Leu Gly Ala Ile Thr Ala Ser Leu Gln Lys Val Glu Asn
      50             55             60

Asp Thr Ser Pro His Arg Glu Arg Ala Thr Leu Leu Glu Glu Gln Leu
      65             70             75             80

Pro Leu Gly Lys Ala Ser Phe His Ile Pro Gln Val Gln Val Arg Asp
      85             90             95

Glu Gly Gln Tyr Gln Cys Ile Ile Ile Tyr Gly Val Ala Trp Asp Tyr
      100             105             110

Lys Tyr Leu Thr Leu Lys Val Lys Ala Ser Tyr Arg Lys Ile Asn Thr
      115             120             125

His Ile Leu Lys Val Pro Glu Thr Asp Glu Val Glu Leu Thr Cys Gln
      130             135             140

Ala Thr Gly Tyr Pro Leu Ala Glu Val Ser Trp Pro Asn Val Ser Val
      145             150             155             160

Pro Ala Asn Thr Ser His Ser Arg Thr Pro Glu Gly Leu Tyr Gln Val
      165             170             175

Thr Ser Val Leu Arg Leu Lys Pro Pro Pro Gly Arg Asn Phe Ser Cys
      180             185             190

Val Phe Trp Asn Thr His Val Arg Glu Leu Thr Leu Ala Ser Ile Asp
      195             200             205

Leu Gln Ser Gln Met Glu Pro Arg Thr His
      210             215

```

<210> 32

<211> 199

<212> PRT

<213> Homo sapiens

<400> 32

```

Leu Phe Thr Val Thr Val Pro Lys Glu Leu Tyr Ile Ile Glu His Gly
 1             5             10             15

Ser Asn Val Thr Leu Glu Cys Asn Phe Asp Thr Gly Ser His Val Asn
      20             25             30

Leu Gly Ala Ile Thr Ala Ser Leu Gln Lys Val Glu Asn Asp Thr Ser
      35             40             45

```

Pro His Arg Glu Arg Ala Thr Leu Leu Glu Glu Gln Leu Pro Leu Gly
 50 55 60
 Lys Ala Ser Phe His Ile Pro Gln Val Gln Val Arg Asp Glu Gly Gln
 65 70 75 80
 Tyr Gln Cys Ile Ile Ile Tyr Gly Val Ala Trp Asp Tyr Lys Tyr Leu
 85 90 95
 Thr Leu Lys Val Lys Ala Ser Tyr Arg Lys Ile Asn Thr His Ile Leu
 100 105 110
 Lys Val Pro Glu Thr Asp Glu Val Glu Leu Thr Cys Gln Ala Thr Gly
 115 120 125
 Tyr Pro Leu Ala Glu Val Ser Trp Pro Asn Val Ser Val Pro Ala Asn
 130 135 140
 Thr Ser His Ser Arg Thr Pro Glu Gly Leu Tyr Gln Val Thr Ser Val
 145 150 155 160
 Leu Arg Leu Lys Pro Pro Pro Gly Arg Asn Phe Ser Cys Val Phe Trp
 165 170 175
 Asn Thr His Val Arg Glu Leu Thr Leu Ala Ser Ile Asp Leu Gln Ser
 180 185 190
 Gln Met Glu Pro Arg Thr His
 195

<210> 33
 <211> 19
 <212> PRT
 <213> Homo sapiens

<400> 33
 Met Ile Phe Leu Leu Leu Met Leu Ser Leu Glu Leu Gln Leu His Gln
 1 5 10 15
 Ile Ala Ala

<210> 34
 <211> 93
 <212> .PRT
 <213> Homo sapiens

<400> 34
 Glu Leu Tyr Ile Ile Glu His Gly Ser Asn Val Thr Leu Glu Cys Asn
 1 5 10 15
 Phe Asp Thr Gly Ser His Val Asn Leu Gly Ala Ile Thr Ala Ser Leu
 20 25 30
 Gln Lys Val Glu Asn Asp Thr Ser Pro His Arg Glu Arg Ala Thr Leu
 35 40 45
 Leu Glu Glu Gln Leu Pro Leu Gly Lys Ala Ser Phe His Ile Pro Gln

50 55 60
 Val Gln Val Arg Asp Glu Gly Gln Tyr Gln Cys Ile Ile Ile Tyr Gly
 65 70 75 80
 Val Ala Trp Asp Tyr Lys Tyr Leu Thr Leu Lys Val Lys
 85 90

 <210> 35
 <211> 94
 <212> PRT
 <213> Homo sapiens

 <400> 35
 Ser Tyr Arg Lys Ile Asn Thr His Ile Leu Lys Val Pro Glu Thr Asp
 1 5 10 15
 Glu Val Glu Leu Thr Cys Gln Ala Thr Gly Tyr Pro Leu Ala Glu Val
 20 25 30
 Ser Trp Pro Asn Val Ser Val Pro Ala Asn Thr Ser His Ser Arg Thr
 35 40 45
 Pro Glu Gly Leu Tyr Gln Val Thr Ser Val Leu Arg Leu Lys Pro Pro
 50 55 60
 Pro Gly Arg Asn Phe Ser Cys Val Phe Trp Asn Thr His Val Arg Glu
 65 70 75 80
 Leu Thr Leu Ala Ser Ile Asp Leu Gln Ser Gln Met Glu Pro
 85 90

 <210> 36
 <211> 301
 <212> PRT
 <213> Homo sapiens

 <400> 36
 Gln Trp Gln Val Phe Gly Pro Asp Lys Pro Val Gln Ala Leu Val Gly
 1 5 10 15
 Glu Asp Ala Ala Phe Ser Cys Phe Leu Ser Pro Lys Thr Asn Ala Glu
 20 25 30
 Ala Met Glu Val Arg Phe Phe Arg Gly Gln Phe Ser Ser Val Val His
 35 40 45
 Leu Tyr Arg Asp Gly Lys Asp Gln Pro Phe Met Gln Met Pro Gln Tyr
 50 55 60
 Gln Gly Arg Thr Lys Leu Val Lys Asp Ser Ile Ala Glu Gly Arg Ile
 65 70 75 80
 Ser Leu Arg Leu Glu Asn Ile Thr Val Leu Asp Ala Gly Leu Tyr Gly
 85 90 95
 Cys Arg Ile Ser Ser Gln Ser Tyr Tyr Gln Lys Ala Ile Trp Glu Leu
 100 105 110

Gln Val Ser Ala Leu Gly Ser Val Pro Leu Ile Ser Ile Ala Gly Tyr
 115 120 125
 Val Asp Arg Asp Ile Gln Leu Leu Cys Gln Ser Ser Gly Trp Phe Pro
 130 135 140
 Arg Pro Thr Ala Lys Trp Lys Gly Pro Gln Gly Gln Asp Leu Ser Thr
 145 150 155 160
 Asp Ser Arg Thr Asn Arg Asp Met His Gly Leu Phe Asp Val Glu Ile
 165 170 175
 Ser Leu Thr Val Gln Glu Asn Ala Gly Ser Ile Ser Cys Ser Met Arg
 180 185 190
 His Ala His Leu Ser Arg Glu Val Glu Ser Arg Val Gln Ile Gly Asp
 195 200 205
 Trp Arg Arg Lys His Gly Gln Ala Gly Lys Arg Lys Tyr Ser Ser Ser
 210 215 220
 His Ile Tyr Asp Ser Phe Pro Ser Leu Ser Phe Met Asp Phe Tyr Ile
 225 230 235 240
 Leu Arg Pro Val Gly Pro Cys Arg Ala Lys Leu Val Met Gly Thr Leu
 245 250 255
 Lys Leu Gln Ile Leu Gly Glu Val His Phe Val Glu Lys Pro His Ser
 260 265 270
 Leu Leu Gln Ile Ser Gly Gly Ser Thr Thr Leu Lys Lys Gly Pro Asn
 275 280 285
 Pro Trp Ser Phe Pro Ser Pro Cys Ala Leu Phe Pro Thr
 290 295 300

<210> 37
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 37
 Met Ala Leu Met Leu Ser Leu Val Leu Ser Leu Leu Lys Leu Gly Ser
 1 5 10 15

Gly

<210> 38
 <211> 26
 <212> PRT
 <213> Homo sapiens

<400> 38
 Thr Ala Ser Pro Trp Met Val Ser Met Thr Val Ile Leu Ala Val Phe
 1 5 10 15

Ile Ile Phe Met Ala Val Ser Ile Cys Cys
 20 25

<210> 39
 <211> 254
 <212> PRT
 <213> Homo sapiens

<400> 39
 Met Glu Pro Ala Ala Leu His Phe Ser Arg Pro Ala Ser Leu Leu
 1 5 10 15
 Leu Leu Leu Ser Leu Cys Ala Leu Val Ser Ala Gln Phe Thr Val Val
 20 25 30
 Gly Pro Ala Asn Pro Ile Leu Ala Met Val Gly Glu Asn Thr Thr Leu
 35 40 45
 Arg Cys His Leu Ser Pro Glu Lys Asn Ala Glu Asp Met Glu Val Arg
 50 55 60
 Trp Phe Arg Ser Gln Phe Ser Pro Ala Val Phe Val Tyr Lys Gly Gly
 65 70 75 80
 Arg Glu Arg Thr Glu Glu Gln Met Glu Glu Tyr Arg Gly Arg Ile Thr
 85 90 95
 Phe Val Ser Lys Asp Ile Asn Arg Gly Ser Val Ala Leu Val Ile His
 100 105 110
 Asn Val Thr Ala Gln Glu Asn Gly Ile Tyr Arg Cys Tyr Phe Gln Glu
 115 120 125
 Gly Arg Ser Tyr Asp Glu Ala Ile Leu Arg Leu Val Val Ala Gly Leu
 130 135 140
 Gly Ser Lys Pro Leu Ile Glu Ile Lys Ala Gln Glu Asp Gly Ser Ile
 145 150 155 160
 Trp Leu Glu Cys Ile Ser Gly Gly Trp Tyr Pro Glu Pro Leu Thr Val
 165 170 175
 Trp Arg Asp Pro Tyr Gly Glu Val Val Pro Ala Leu Lys Glu Val Ser
 180 185 190
 Ile Ala Asp Ala Asp Gly Leu Phe Met Val Thr Thr Ala Val Ile Ile
 195 200 205
 Arg Asp Lys Tyr Val Arg Asn Val Ser Cys Ser Val Asn Asn Thr Leu
 210 215 220
 Leu Gly Gln Glu Lys Glu Thr Val Ile Phe Ile Pro Glu Ser Phe Met
 225 230 235 240
 Pro Ser Ala Ser Pro Trp Met Val Ala Leu Ala Val Ile Leu
 245 250

<210> 40
 <211> 227
 <212> PRT
 <213> Homo sapiens

<400> 40

Gln Phe Thr Val Val Gly Pro Ala Asn Pro Ile Leu Ala Met Val Gly
 1 5 10 15

Glu Asn Thr Thr Leu Arg Cys His Leu Ser Pro Glu Lys Asn Ala Glu
 20 25 30

Asp Met Glu Val Arg Trp Phe Arg Ser Gln Phe Ser Pro Ala Val Phe
 35 40 45

Val Tyr Lys Gly Gly Arg Glu Arg Thr Glu Glu Gln Met Glu Glu Tyr
 50 55 60

Arg Gly Arg Ile Thr Phe Val Ser Lys Asp Ile Asn Arg Gly Ser Val
 65 70 75 80

Ala Leu Val Ile His Asn Val Thr Ala Gln Glu Asn Gly Ile Tyr Arg
 85 90 95

Cys Tyr Phe Gln Glu Gly Arg Ser Tyr Asp Glu Ala Ile Leu Arg Leu
 100 105 110

Val Val Ala Gly Leu Gly Ser Lys Pro Leu Ile Glu Ile Lys Ala Gln
 115 120 125

Glu Asp Gly Ser Ile Trp Leu Glu Cys Ile Ser Gly Gly Trp Tyr Pro
 130 135 140

Glu Pro Leu Thr Val Trp Arg Asp Pro Tyr Gly Glu Val Val Pro Ala
 145 150 155 160

Leu Lys Glu Val Ser Ile Ala Asp Ala Asp Gly Leu Phe Met Val Thr
 165 170 175

Thr Ala Val Ile Ile Arg Asp Lys Tyr Val Arg Asn Val Ser Cys Ser
 180 185 190

Val Asn Asn Thr Leu Leu Gly Gln Glu Lys Glu Thr Val Ile Phe Ile
 195 200 205

Pro Glu Ser Phe Met Pro Ser Ala Ser Pro Trp Met Val Ala Leu Ala
 210 215 220

Val Ile Leu
 225

<210> 41

<211> 27

<212> PRT

<213> Homo sapiens

<400> 41

Met Glu Pro Ala Ala Ala Leu His Phe Ser Arg Pro Ala Ser Leu Leu
 1 5 10 15

Leu Leu Leu Ser Leu Cys Ala Leu Val Ser Ala
 20 25

<210> 42
 <211> 20
 <212> PRT
 <213> Homo sapiens

<400> 42
 Gly Pro Thr Gly Ala Arg Leu Thr Leu Val Leu Ala Leu Thr Val Ile
 1 5 10 15
 Leu Glu Leu Thr
 20

<210> 43
 <211> 394
 <212> PRT
 <213> Homo sapiens

<400> 43
 Met Arg Glu Ile Val Trp Tyr Arg Val Thr Asp Gly Gly Thr Ile Lys
 1 5 10 15
 Gln Lys Ile Phe Thr Phe Asp Ala Met Phe Ser Thr Asn Tyr Ser His
 20 25 30
 Met Glu Asn Tyr Arg Lys Arg Glu Asp Leu Val Tyr Gln Ser Thr Val
 35 40 45
 Arg Leu Pro Glu Val Arg Ile Ser Asp Asn Gly Pro Tyr Glu Cys His
 50 55 60
 Val Gly Ile Tyr Asp Arg Ala Thr Arg Glu Lys Val Val Leu Ala Ser
 65 70 75 80
 Gly Asn Ile Phe Leu Asn Val Met Ala Pro Pro Thr Ser Ile Glu Val
 85 90 95
 Val Ala Ala Asp Thr Pro Ala Pro Phe Ser Arg Tyr Gln Ala Gln Asn
 100 105 110
 Phe Thr Leu Val Cys Ile Val Ser Gly Gly Lys Pro Ala Pro Met Val
 115 120 125
 Tyr Phe Lys Arg Asp Gly Glu Pro Ile Asp Ala Val Pro Leu Ser Glu
 130 135 140
 Pro Pro Ala Ala Ser Ser Gly Pro Leu Gln Asp Ser Arg Pro Phe Arg
 145 150 155 160
 Ser Leu Leu His Arg Asp Leu Asp Asp Thr Lys Met Gln Lys Ser Leu
 165 170 175
 Ser Leu Leu Asp Ala Glu Asn Arg Gly Gly Arg Pro Tyr Thr Glu Arg
 180 185 190
 Pro Ser Arg Gly Leu Thr Pro Asp Pro Asn Ile Leu Leu Gln Pro Thr
 195 200 205
 Thr Glu Asn Ile Pro Glu Thr Val Val Ser Arg Glu Phe Pro Arg Trp
 210 215 220

Val His Ser Ala Glu Pro Thr Tyr Phe Leu Arg His Ser Arg Thr Pro
 225 230 235 240
 Ser Ser Asp Gly Thr Val Glu Val Arg Ala Leu Leu Thr Trp Thr Leu
 245 250 255
 Asn Pro Gln Ile Asp Asn Glu Ala Leu Phe Ser Cys Glu Val Lys His
 260 265 270
 Pro Ala Leu Ser Met Pro Met Gln Ala Glu Val Thr Leu Val Ala Pro
 275 280 285
 Lys Gly Pro Lys Ile Val Met Thr Pro Ser Arg Ala Arg Val Gly Asp
 290 295 300
 Thr Val Arg Ile Leu Val His Gly Phe Gln Asn Glu Val Phe Pro Glu
 305 310 315 320
 Pro Met Phe Thr Trp Thr Arg Val Gly Ser Arg Leu Leu Asp Gly Ser
 325 330 335
 Ala Glu Phe Asp Gly Lys Glu Leu Val Leu Glu Arg Val Pro Ala Glu
 340 345 350
 Leu Asn Gly Ser Met Tyr Arg Cys Thr Ala Gln Asn Pro Leu Gly Ser
 355 360 365
 Thr Asp Thr His Thr Arg Leu Ile Val Phe Glu Asn Pro Asn Ile Pro
 370 375 380
 Arg Gly Thr Glu Asp Ser Asn Gly Ser Ile
 385 390

<210> 44
 <211> 132
 <212> PRT
 <213> Homo sapiens

<400> 44
 Gln Val Thr Val Val Gly Pro Thr Asp Pro Ile Leu Ala Met Val Gly
 1 5 10 15
 Glu Asn Thr Thr Leu Arg Cys Cys Leu Ser Pro Glu Glu Asn Ala Glu
 20 25 30
 Asp Met Glu Val Arg Trp Phe Gln Ser Gln Phe Ser Pro Ala Val Phe
 35 40 45
 Val Tyr Lys Gly Gly Arg Glu Arg Thr Glu Glu Gln Lys Glu Glu Tyr
 50 55 60
 Arg Gly Arg Thr Thr Phe Val Ser Lys Asp Ser Arg Gly Ser Val Ala
 65 70 75 80
 Leu Ile Ile His Asn Val Thr Ala Glu Asp Asn Gly Ile Tyr Gln Cys
 85 90 95
 Tyr Phe Gln Glu Gly Arg Ser Cys Asn Glu Ala Ile Leu His Leu Val
 100 105 110

Val Ala Asp Gln His Asn Pro Leu Ser Trp Ile Pro Ile Pro Gln Gly
 115 120 125

Thr Leu Ser Leu
 130

<210> 45
 <211> 27
 <212> PRT
 <213> Homo sapiens

<400> 45
 Met Glu Pro Ala Ala Ala Leu His Phe Ser Arg Pro Ala Ser Leu Leu
 1 5 10 15

Leu Leu Leu Ser Leu Cys Ala Leu Val Ser Ala
 20 25

<210> 46
 <211> 13
 <212> PRT
 <213> Homo sapiens

<400> 46
 Leu Gly Ile Leu Cys Cys Gly Leu Phe Phe Gly Ile Val
 1 5 10

<210> 47
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 47
 Met Ala Leu Met Leu Ser Leu Val Leu Ser Leu Leu Lys Leu Gly Ser
 1 5 10 15

Gly

<210> 48
 <211> 239
 <212> PRT
 <213> Homo sapiens

<400> 48
 Met Ala Leu Met Leu Ser Leu Val Leu Ser Leu Leu Lys Leu Gly Ser
 1 5 10 15

Gly Gln Trp Gln Val Phe Gly Pro Asp Lys Pro Val Gln Ala Leu Val
 20 25 30

Gly Glu Asp Ala Ala Phe Ser Cys Phe Leu Ser Pro Lys Thr Asn Ala
 35 40 45

Glu Ala Met Glu Val Arg Phe Phe Arg Gly Gln Phe Ser Ser Val Val
 50 55 60

His Leu Tyr Arg Asp Gly Lys Asp Gln Pro Phe Met Gln Met Pro Gln
 65 70 75 80
 Tyr Gln Gly Arg Thr Lys Leu Val Lys Asp Ser Ile Ala Glu Gly Arg
 85 90 95
 Ile Ser Leu Arg Leu Glu Asn Ile Thr Val Leu Asp Ala Gly Leu Tyr
 100 105 110
 Gly Cys Arg Ile Ser Ser Gln Ser Tyr Tyr Gln Lys Ala Ile Trp Glu
 115 120 125
 Leu Gln Val Ser Ala Leu Gly Ser Val Pro Leu Ile Ser Ile Thr Gly
 130 135 140
 Tyr Val Asp Arg Asp Ile Gln Leu Leu Cys Gln Ser Ser Gly Trp Phe
 145 150 155 160
 Pro Arg Pro Thr Ala Lys Trp Lys Gly Pro Gln Gly Gln Asp Leu Ser
 165 170 175
 Thr Asp Ser Arg Thr Asn Arg Asp Met His Gly Leu Phe Asp Val Glu
 180 185 190
 Ile Ser Leu Thr Val Gln Glu Asn Ala Gly Ser Ile Ser Cys Ser Met
 195 200 205
 Arg His Ala His Leu Ser Arg Glu Val Glu Ser Arg Val Gln Ile Gly
 210 215 220
 Asp Thr Phe Phe Glu Pro Ile Ser Trp His Leu Ala Thr Lys Val
 225 230 235

<210> 49

<211> 222

<212> PRT

<213> Homo sapiens

<400> 49

Gln Trp Gln Val Phe Gly Pro Asp Lys Pro Val Gln Ala Leu Val Gly
 1 5 10 15
 Glu Asp Ala Ala Phe Ser Cys Phe Leu Ser Pro Lys Thr Asn Ala Glu
 20 25 30
 Ala Met Glu Val Arg Phe Phe Arg Gly Gln Phe Ser Ser Val Val His
 35 40 45
 Leu Tyr Arg Asp Gly Lys Asp Gln Pro Phe Met Gln Met Pro Gln Tyr
 50 55 60
 Gln Gly Arg Thr Lys Leu Val Lys Asp Ser Ile Ala Glu Gly Arg Ile
 65 70 75 80
 Ser Leu Arg Leu Glu Asn Ile Thr Val Leu Asp Ala Gly Leu Tyr Gly
 85 90 95
 Cys Arg Ile Ser Ser Gln Ser Tyr Tyr Gln Lys Ala Ile Trp Glu Leu
 100 105 110

Gln Val Ser Ala Leu Gly Ser Val Pro Leu Ile Ser Ile Thr Gly Tyr
115 120 125

Val Asp Arg Asp Ile Gln Leu Leu Cys Gln Ser Ser Gly Trp Phe Pro
130 135 140

Arg Pro Thr Ala Lys Trp Lys Gly Pro Gln Gly Gln Asp Leu Ser Thr
145 150 155 160

Asp Ser Arg Thr Asn Arg Asp Met His Gly Leu Phe Asp Val Glu Ile
165 170 175

Ser Leu Thr Val Gln Glu Asn Ala Gly Ser Ile Ser Cys Ser Met Arg
180 185 190

His Ala His Leu Ser Arg Glu Val Glu Ser Arg Val Gln Ile Gly Asp
195 200 205

Thr Phe Phe Glu Pro Ile Ser Trp His Leu Ala Thr Lys Val
210 215 220

**INDICATIONS RELATING TO A DEPOSITED MICROORGANISM
OR OTHER BIOLOGICAL MATERIAL**

(PCT Rule 13bis)

A. The indications made below relate to the deposited microorganism or other biological material referred to in the description at Page 115, Table 1.

B. IDENTIFICATION OF DEPOSIT

Further deposits are identified on an additional sheet ☐

Name of depositary institution: American Type Culture Collection

Address of depositary institution (including postal code and country)

10801 University Boulevard
Manassas, Virginia 20110-2209
United States of America

Date of deposit

August 7, 2000

Accession Number

PTA-2332

C. ADDITIONAL INDICATIONS (leave blank if not applicable)

This information is continued on an additional sheet ☐

D. DESIGNATED STATES FOR WHICH INDICATIONS ARE MADE (if the indications are not for all designated States)

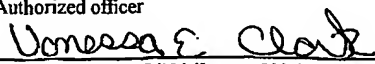
Europe

In respect of those designations in which a European Patent is sought a sample of the deposited microorganism will be made available until the publication of the mention of the grant of the European patent or until the date on which the application has been refused or withdrawn or is deemed to be withdrawn, only by the issue of such a sample to an expert nominated by the person requesting the sample (Rule 28(4) EPC).

Continued on additional sheets

E. SEPARATE FURNISHING OF INDICATIONS (leave blank if not applicable)

The indications listed below will be submitted to the international Bureau later (specify the general nature of the indications e.g., "Accession Number of Deposit")

	For receiving Office use only			For International Bureau use only	
<input checked="" type="checkbox"/> This sheet was received with the international application			<input type="checkbox"/> This sheet was received by the International Bureau on:		
Authorized officer 			Authorized officer		

ATCC Deposit No. PTA-2332

CANADA

The applicant requests that, until either a Canadian patent has been issued on the basis of an application or the application has been refused, or is abandoned and no longer subject to reinstatement, or is withdrawn, the Commissioner of Patents only authorizes the furnishing of a sample of the deposited biological material referred to in the application to an independent expert nominated by the Commissioner, the applicant must, by a written statement, inform the International Bureau accordingly before completion of technical preparations for publication of the international application.

NORWAY

The applicant hereby requests that the application has been laid open to public inspection (by the Norwegian Patent Office), or has been finally decided upon by the Norwegian Patent Office without having been laid open inspection, the furnishing of a sample shall only be effected to an expert in the art. The request to this effect shall be filed by the applicant with the Norwegian Patent Office not later than at the time when the application is made available to the public under Sections 22 and 33(3) of the Norwegian Patents Act. If such a request has been filed by the applicant, any request made by a third party for the furnishing of a sample shall indicate the expert to be used. That expert may be any person entered on the list of recognized experts drawn up by the Norwegian Patent Office or any person approved by the applicant in the individual case.

AUSTRALIA

The applicant hereby gives notice that the furnishing of a sample of a microorganism shall only be effected prior to the grant of a patent, or prior to the lapsing, refusal or withdrawal of the application, to a person who is a skilled addressee without an interest in the invention (Regulation 3.25(3) of the Australian Patents Regulations).

FINLAND

The applicant hereby requests that, until the application has been laid open to public inspection (by the National Board of Patents and Regulations), or has been finally decided upon by the National Board of Patents and Registration without having been laid open to public inspection, the furnishing of a sample shall only be effected to an expert in the art.

ATCC Deposit No. PTA-2332

UNITED KINGDOM

The applicant hereby requests that the furnishing of a sample of a microorganism shall only be made available to an expert. The request to this effect must be filed by the applicant with the International Bureau before the completion of the technical preparations for the international publication of the application.

DENMARK

The applicant hereby requests that, until the application has been laid open to public inspection (by the Danish Patent Office), or has been finally decided upon by the Danish Patent office without having been laid open to public inspection, the furnishing of a sample shall only be effected to an expert in the art. The request to this effect shall be filed by the applicant with the Danish Patent Office not later than at the time when the application is made available to the public under Sections 22 and 33(3) of the Danish Patents Act. If such a request has been filed by the applicant, any request made by a third party for the furnishing of a sample shall indicate the expert to be used. That expert may be any person entered on a list of recognized experts drawn up by the Danish Patent Office or any person by the applicant in the individual case.

SWEDEN

The applicant hereby requests that, until the application has been laid open to public inspection (by the Swedish Patent Office), or has been finally decided upon by the Swedish Patent Office without having been laid open to public inspection, the furnishing of a sample shall only be effected to an expert in the art. The request to this effect shall be filed by the applicant with the International Bureau before the expiration of 16 months from the priority date (preferably on the Form PCT/RO/134 reproduced in annex Z of Volume I of the PCT Applicant's Guide). If such a request has been filed by the applicant any request made by a third party for the furnishing of a sample shall indicate the expert to be used. That expert may be any person entered on a list of recognized experts drawn up by the Swedish Patent Office or any person approved by a applicant in the individual case.

NETHERLANDS

The applicant hereby requests that until the date of a grant of a Netherlands patent or until the date on which the application is refused or withdrawn or lapsed, the microorganism shall be made available as provided in the 31F(1) of the Patent Rules only by the issue of a sample to an expert. The request to this effect must be furnished by the applicant with the Netherlands Industrial Property Office before the date on which the application is made available to the public under Section 22C or Section 25 of the Patents Act of the Kingdom of the Netherlands, whichever of the two dates occurs earlier.